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IZABELA KAŁUCKA

**MACROFUNGI IN THE SECONDARY SUCCESSION  
ON THE ABANDONED FARMLAND  
NEAR THE BIAŁOWIEŻA OLD-GROWTH FOREST**

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Department of Algology and Mycology  
University of Łódź  
12/16 Banacha Str.  
PL-90-237 Łódź

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## CONTENTS

1. Introduction .....	5
1.1. Fungal succession and old-fields .....	5
1.2. Scheme of spontaneous secondary succession on abandoned fields near the Białowieża old-growth forest .....	8
2. Subject and aims of study .....	12
3. Study area .....	13
4. Material and methods .....	15
5. Results .....	25
5.1. Fungi in permanent plots .....	25
5.2. Changes in species and sporocarp distribution along successional gradient .....	34
5.2.1. Species richness and diversity .....	34
5.2.2. Species abundance patterns .....	39
5.2.3. Spatial frequency patterns .....	53
5.2.4. Ectomycorrhizal carpophore production .....	53
5.3. Similarity structure of macrofungi .....	56
5.4. Quantitative changes in ectomycorrhizal roots along successional gradient .....	60
6. Discussion .....	61
6.1. Macrofungi in the scheme of spontaneous secondary succession on the abandoned fields near Białowieża .....	62
6.2. Patterns of fungal succession in post-agricultural land .....	66
7. Summary of results and conclusions .....	81
8. References .....	85
Appendix 1. Soil analysis in the permanent plots .....	95
Appendix 2. Vegetation description in the permanent plots .....	101
Appendix 3. Macrofungi in the plots representing subsequent phases of vegetation succession in the old-fields near the Białowieża old-growth forest .....	115
Appendix 4. Results of k-mean analysis of fungi .....	141
Appendix 5. List of Basidiomycota in the old-field successional series near the Białowieża old-growth forest (Jelonka reserve and Czechy Orłańskie reserve) with notes on distribution and substrates .....	145

## ABSTRACT

Izabela KAŁUCKA. *Macrofungi in the secondary succession on the abandoned farmland near the Białowieża old-growth forest*. Monogr. Bot., Vol. 99, 155 pp., 2009.

The paper deals with the succession of basidiomycete macrofungi on abandoned farmland. The research was carried out in the unique complex of old fields of different age located on the southwest edge of the Białowieża old-growth forest (NE Poland). These lands undergo spontaneous secondary succession leading to formation of a continental fresh pine forest. Regular observations of basidiomata in 16 permanent plots (11 plots of 1000 m<sup>2</sup>, 5 plots of 400 m<sup>2</sup>) representing initial, optimal and terminal stage of vegetation development from psammophilous grassland to a pine forest with 100-year-old tree stand were conducted for three years. Soil and vegetation analyses were also carried out as well as quantitative examination of ectomycorrhizae in root samples.

The series yielded 300 species of macromycetes (including 6 collective taxa) found in permanent plots plus 18 species outside the plots. The species composition, spatial distribution of sporocarps, and quantitative structure of fungal communities changed along the successional gradient. Increasing diversity of fungi resulted from persistence of species typical of open areas and species associated with younger trees, as well as accumulation of species associated with older trees and more developed stands. Some species typical of younger phases were replaced with species arriving later in the chronosequence. Species usually occurred long before and after attaining maximum abundance and frequency. Species associated with certain periods of vegetation development could be distinguished, as well as species accompanying the phytocoenoses regardless of their developmental stage. Patterns in structural changes of the fungal community, e.g., in number of species, abundance, frequency and production of carpophores, could also be recognised. They were different in ectomycorrhizal and in saprotrophic fungi. The occurrence of the former was influenced mainly by the presence of symbiotic trees and their age, stand structure and soil properties, while occurrence of the latter was affected by microclimatic factors and by diversity of accessible substrates. Changes in fungal communities were consistent with vegetation changes, but were also specific to this group of organisms.

*Key words: old-field succession, fungal succession, ectomycorrhizal fungi, saprotrophic fungi, sporocarp production, NE Poland*



## 1. INTRODUCTION

The influence of fungi on the formation and functioning of ecosystems is more complex and diverse than usually appreciated. Fungi take part in all major ecological processes, either in direct or indirect ways, and the existence of a great variety of other organisms and whole ecosystems depends on their activity. As decomposers, they drive the global carbon cycle by preventing an excessive accumulation of organic matter and its breakdown into the compounds which can be utilized by other organisms. As mycorrhizal partners fungi enhance plants water uptake and mineral nutrition, they supply regulatory substances and protect against pathogens and environmental stress. They interconnect individual plants via common mycorrhizal networks and build extensive sink-source relationships affecting the process of primary production as well as the composition and dynamics of vegetation. Fungal endophytes inhabiting tissues of most plants can considerably influence their condition and alter adaptation capabilities. As lichen compounds, fungi contribute to the colonization of most pioneer habitats on earth and they provide a shelter for the important nitrogen-fixing organisms – cyanobacteria. As parasites they strongly influence populations of their hosts and act as biological control. Also, fungi serve as a habitat or food for different organisms including bacteria, other fungi, nematodes, insects and mammals (RAYNER 1992; TRAPPE & LUOMA 1992; DIGHTON 2003). According to RAYNER (1992) *In short, fungi demand the attention of anyone eager to understand the complex interactions and feedback governing life processes in the biosphere.*

### 1.1. Fungal succession and old-fields

One of the key life processes is succession. It is a process of initiation and development of a community (and whole biocoenosis) within the confines of a potential bioclimate and relatively uniform environmental conditions (FALIŃSKI 1998). Succession of both plants and fungi can be defined as a directional change in the composition, relative abundance and spatial pattern of species comprising communities (FRANKLAND 1992, 1998). In strict mycological sense it is a sequential occupation of the same site by thalli (normally mycelia), either of different fungi, or of different associations of fungi (RAYNER & TODD 1979). The term fungal community is frequently used in this context meaning assemblage of fungi living together in the same time and space (BEGON *et al.* 1996) or occurring within a certain uniform habitat (ARNOLDS 1992a).

Two different types of fungal succession can be distinguished depending on the type of resource used (PARK 1968; FRANKLAND 1992): substratum succession and seral succession. The former means time-related changes in the community structure of fungi bound to the substrate usually limited in space which offers unrenovable resource, such as fungi inhabiting litter or wood (e.g., DICKINSON & PUGH 1974; RAYNER & TODD 1979). The latter refers to a series of fungal assemblages following successional changes in vegetation where the resource in the short-term is renewable, for example mycorrhizal fungi associated

with their host plants (e.g., ALLEN E.B. & ALLEN M.F. 1990; HELM *et al.* 1996; NARA *et al.* 2003a).

Chronosequences of fungal communities occur both in primary and secondary successional systems. Primary succession involves the colonization and establishment of vegetation on newly exposed substrates, usually scarce in nutrients, barren and devoid of autochthonous organic matter, in natural as well as anthropogenic sites. Successional series of ectomycorrhizal (ECM) fungi have been described in glacier forefronts (e.g., HELM *et al.* 1996; CÁZARES *et al.* 2005; MÜHLMANN & PEINTNER 2008), volcanic deserts (e.g., ALLEN M.F. *et al.* 1992; NARA *et al.* 2003a, b; OBASE *et al.* 2007), in coastal dunes (e.g., DOMINIK 1951; DOMINIK & PACHLEWSKI 1955; ALLEN M.F. & ALLEN E.B. 1992; VAN DER HEIJDEN *et al.* 1999; VAN DER HEIJDEN & VOSATKA 1999; ASHKANNEJHAD & HORTON 2006), coal mine spoil (e.g., LISIEWSKA & SIEDLACZEK 1982; DANIELSON 1991; GEBHARDT *et al.* 2007), ski slopes and road construction sites (e.g., HASHIMOTO & HYAKUMACHI 2000). Secondary succession involves the replacement of pre-existing vegetation on developed soil containing the residues left by the previous community as a result of disturbance, for instance burned areas or on abandoned fields. Changes in fungal communities in different post-fire ecosystems have been described, e.g., by VISSER (1995), HORTON *et al.* (1998), FRIEDRICH (2001), SUMOROK (2001). Fungal dynamics in post-agricultural areas have also been extensively studied, however most of the research deals with soil microfungi or arbuscular mycorrhizae (AM) (JOHNSON *et al.* 1991; BOERNER *et al.* 1996; BARNI & SINISCALCO 2000; HEDLUND 2002; HEDLUND & GORMSEN 2002; RICHTER *et al.* 2002; KLAMER & HEDLUND 2004). It has been shown that agricultural management can reduce the abundance and diversity of these organisms, hence the rate of recovery of the vegetation and post-disturbance plant species composition can be affected (REEVES *et al.* 1979; VAN DER HEIJDEN *et al.* 1998).

The ECM inoculum potential and colonization in post-arable fields have been less studied, despite being crucial for afforestation practices (PERRY *et al.* 1987; KROPP & LANGLOIS 1990; LE TACON *et al.* 1994). BOERNER *et al.* (1996) compared an active agricultural field, 5- and 10-year-old fields, and undisturbed mature forest. They found that in the active field, ECM inoculum was sparse and patchily distributed, surrounded by large areas without ECM inoculum, and correlated with the ploughing pattern, so the chance of any seeds of ECM-dependent plants to find mycorrhizal partners (and thus effectively colonize the site) was very low. Both the ECM infectiveness and its homogeneity, and the ECM diversity increased with time since last disturbance, reaching the state of a *virtually unbroken matrix of high ECM infectiveness* in mature forest. However, the data suggested that the pre-disturbance ECM fungi level could not be re-established in less than 25-30 years following agriculture cessation.

HEDLUND & GORMSEN (2002) revealed that ECM colonization and the survival of tree seedlings in abandoned fields were higher in areas that were naturally colonized by plants and not managed by sowing additional species. The effect of artificial inoculation of pine and spruce seedlings outplanted into a field abandoned for more than 10 years has been studied by MENKIS *et al.* (2007). It appeared that even after the first year natural mycorrhizal colonization prevailed in the preinoculated root systems and introduced mycorrhizas were seldom found. The impact of inoculation on subsequent ECM communities was low and after two growing seasons extensive natural mycorrhization took place resulting in quite diverse fungal communities. This indicated that ECM formation in root systems is governed mainly by environmental factors, at least in not harsh conditions, and after sufficient passage of time from abandonment. However, the tree seedlings inoculated

with *Cenococcum geophilum* and *Piceirhiza bicolorata* (but not *Hebeloma crustuliniforme*) showed significantly higher survival and growth when compared with controls over a longer period.

An interesting field experiment examining the influence of distance from established vegetation on the presence, abundance, diversity and community composition of ECM fungi in abandoned agricultural fields has been carried out by DICKIE & REICH (2005). Distinct spatial complexity of these fungi can strongly influence the establishment of tree seedlings and thus development of forest-type vegetation. The ECM fungal community varied considerably with distance from the forest edge. The abundance of mycorrhizae, uniformly high near trees, declined rapidly around 15 m from the base of trees. Similar decrease was observed in ECM species richness.

This kind of research focused on anthropogenic as well as natural factors influencing plant colonization patterns of post-agricultural land, especially those leading to successful establishment of trees and forest development, is of particular importance. This is particularly true when serious difficulties arise, as frequently happens (DOMINIK 1961; PERRY *et al.* 1987; LEPŠ & REJMANEK 1991; SOB CZAK 1990; SZUJECKI 1990; SIEROTA 1996). Also, the rates of secondary succession of such habitats and recovery to late successional woody vegetation vary considerably, from 10-40 years (KEEVER 1983) to 140 years (European mesophilous pine forest) and 350 years (European mixed hardwood forest) (FALIŃSKI 1986a). In the Alps, 100 years after pasture abandonment, some areas are still in an early stage of transition from grassland to savannah-like community (WIEMKEN & BOLLER 2006) and delayed succession is ascribed to the limitations connected with formation of ectomycorrhizae.

Post-agricultural areas proved to be good model systems for the studies of ECM fungal chronosequences following disturbance and for general fungal ecology.

The examination of ECM fungi sequences in young plantations of birch, *Betula pendula* and *B. pubescens* (MASON *et al.* 1982, 1983) on post-agricultural land gave rise to the successional concept of 'early- and late-stage fungi'. The results, confirmed in other types of forests (CHU-CHOU & GRACE 1981, 1988; DIGHTON *et al.* 1986; NATARAJAN *et al.* 1992; RICHTER & BRUHN 1993) showed that species of fungi occurring at the early stages of woodland development ('early-stage fungi', e.g., species of *Hebeloma*, *Inocybe*, *Laccaria*, *Suillus* and *Rhizopogon*) are, in due course, superseded or joined by others ('late-stage fungi', e.g., species of *Russula*, *Cortinarius*, *Amanita*, *Lactarius*, *Leccinum*). The concept was also discussed in the context of life histories of the species (DIGHTON & MASON 1985; ANDREWS 1992). The composition of the ECM fungal community is determined by r-selection at the beginning of the successional process, while K-selection prevails when trees age and organic matter accumulates. The early-stage fungi, colonizing seedlings, have a low carbohydrate demand (supplied by the host tree), they exploit nutrients contained in the inorganic pool, are capable of rapid mycelial growth and spread mainly by spores. The late-stage fungi are associated with older/larger trees or older forests, have a high carbohydrate demand, derive nutrients mainly from the organic pool, exhibit high competitive abilities due to the production of mycelial strands, and spread mainly via vegetative mycelium.

An ordered chronosequence of the ECM fungi was first ascribed to the age of host trees and the temporal stage of woodland development, and to changes in host physiology, photosynthate allocation in particular, as a result of aging (DIGHTON & MASON 1985). However, it was also found that soil conditions, accumulation of litter, development of humus layer and content of organic matter are of great importance, as well as the stand structure (e.g., LAST *et al.* 1987; BLASIUS & OBERWINKLER 1989; DE VRIES *et al.* 1995; JUMPPONEN *et al.* 1999).

## 1.2. Scheme of spontaneous secondary succession on abandoned fields near the Białowieża old-growth forest

Successional processes in nature are usually long-lasting and slow, thus the examinations of the successional series developing in one place are usually carried out, with caution in the interpretation of the results, with the method of “time for space replacement” (JOHNSON & MIYANISHI 2008). A rare example of a site which is suitable for this kind of research is a complex of old fields situated near the southwest peripheries of the Białowieża National Park in the Białowieża old-growth forest (Fig.1), one of the best preserved primeval forests in Europe, which is listed by UNESCO as a World Heritage Site. In the past, this area lay in the borderland between two huge forest complexes – Białowieża and Bielsk old-growth forest, the latter not having existed for five hundred

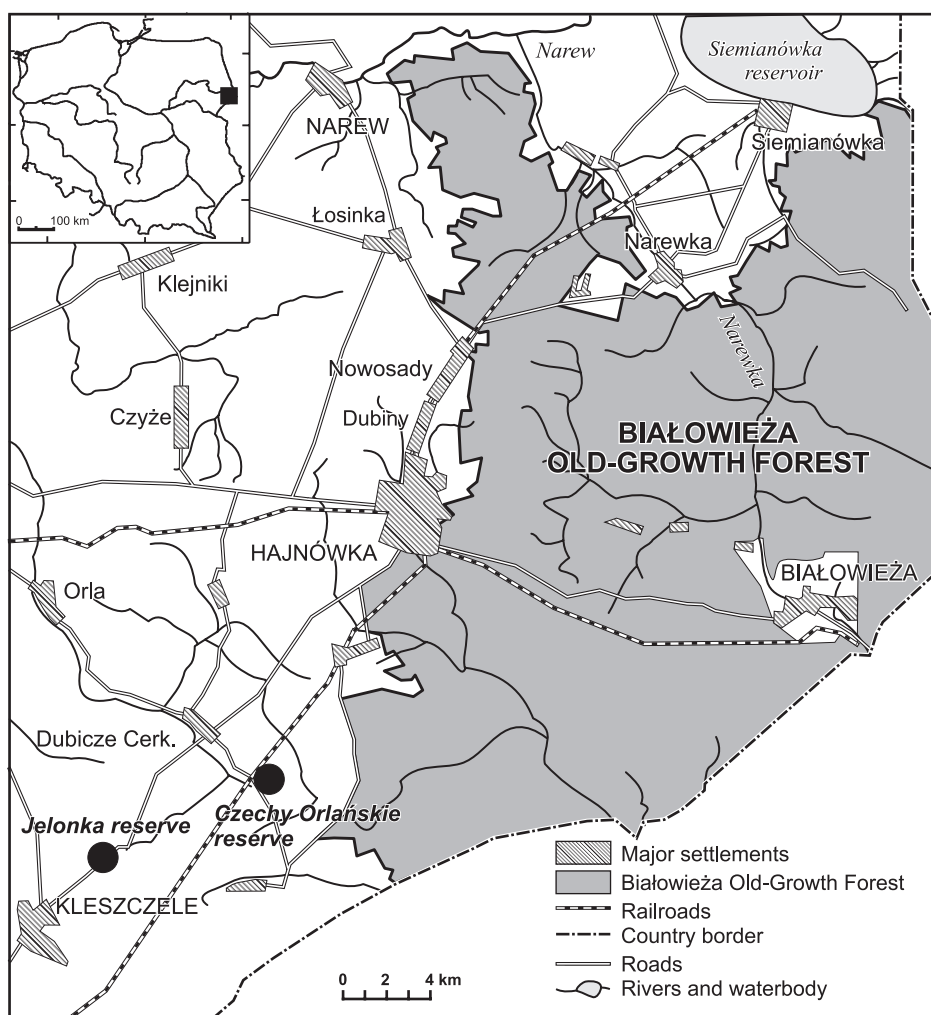


Fig. 1. The study area – Jelonka and Czechy Orlańskie reserves.

years (FALIŃSKI 1986b). As a result of growing settlement and farming it became devoid of forest cover in the XVI century (ALEXANDROWICZ 1964; WIŚNIEWSKI 1964). However, as the soil here is poor and was soon depleted of its nutrients, part of the area was gradually left aside and spontaneously afforested again (KONDRATIUK 1974). The process intensified following the I and II World Wars. The main body of the abandoned fields arose as a result of demographic changes and the changes in agrarian policy in the last 50-60 years on rather dry and sandy soils which potentially are the habitat of mesophilous (fresh) pine forest. The area was subject to the government programme of land reclamation (up to 1992 ca 12,000 ha were afforested; FALIŃSKI *et al.* 1993). The most valuable part was excluded from management and secured for ecological studies as a nature reserve, which makes it unique on a European scale.

There occur side by side strips of old fields with vegetation communities representing initial, optimal and terminal stage of succession series from psammophilous grassland to continental fresh pine forest *Peucedano-Pinetum*. As a result of long-term studies initiated in 1970, ten phases of vegetation development were distinguished, as well as their duration (based on dendrochronometric studies) and dynamics of main components – vascular plants, bryophytes and lichens (Fig. 2) (FALIŃSKI *et al.* 1993; FALIŃSKI 1998, 2002). The syndynamic research was carried out mainly in the Jelonka reserve; the pine forest representing nearly mature final vegetation community, growing in the Czechy Orlańskie reserve, was included. A detailed description of the process of secondary succession, with diagnoses of consecutive phases of vegetation development, was published by FALIŃSKI (1986b, 1998) and FALIŃSKI *et al.* (1993). In brief, they are as follows:

**Phase 0. A segetal community.** *Teesdaleo-Arno-seridetum* Malc. 1929 R. Tx. 1937 [= *Arno-serido-Scleranthetum* (Chouard 1925) R. Tx. 1937] – a weed association poor in species, accompanying the crops of rye, oats, potato, yellow lupine and buckwheat. Among frequent species are *Rumex acetosella*, *Scleranthus annuus*, *S. perennis*, *Convolvulus arvensis*, *Viola arvensis*, *Sinapis arvensis*, *Arabidopsis thaliana*, *Festuca rubra*, *Anthemis arvensis*, *Stachys annua* and *Galeopsis ladanum*. Some strongly invasive pioneer species of mosses can appear, like *Ceratodon purpureus* and *Brachythecium albicans*.

**Phase 1. The pioneer phase of a psammophilous herb-grassland on a new fallow.** Duration 1-2 years after the field abandonment. Some typical weeds still occur, *Galeopsis ladanum*, *Stachys annua*, *Digitaria ischaemum*, *Spergula arvensis* and *Gypsophila muralis*. They are accompanied by herbaceous perennials, among which typical psammophytes prevail: *Jasione montana*, *Helichrysum arenarium*, *Artemisia campestris*, *Hypericum perforatum*, *Solidago virgaurea*, *Hypochoeris radicata*, *Rumex acetosella*, *Hieracium pilosella*, *Knautia arvensis* and *Achillea millefolium*. The occurrence of psammophilous grass *Corynephorus canescens* as well as therophytes like *Filago minima*, *Gnaphalium sylvaticum* and *Scleranthus annuus* is also characteristic. The vegetation community is a psammophilous herb-grassland belonging to the class *Koelerietea*, and the order of *Festuco-Sedetalia*. It represents a type of inland grassland of continental character, belonging to the alliance *Koelerion glaucae* (Volk 1931) Klika 1935 and the community *Koelerio-Astragaletum arenariae* described by GŁOWACKI (1988). The optimum of this community occurs in phases 4-6. As early as during phase 1 pioneer species of mosses can appear, *Polytrichum piliferum* and *Niphotrichum canescens*.

**Phases 2, 3 and 4. A psammophilous herb-grassland with lichen communities.** Duration 2-15 years after the field abandonment. *Thymus serpyllum* and *Astragalus arenarius* appear



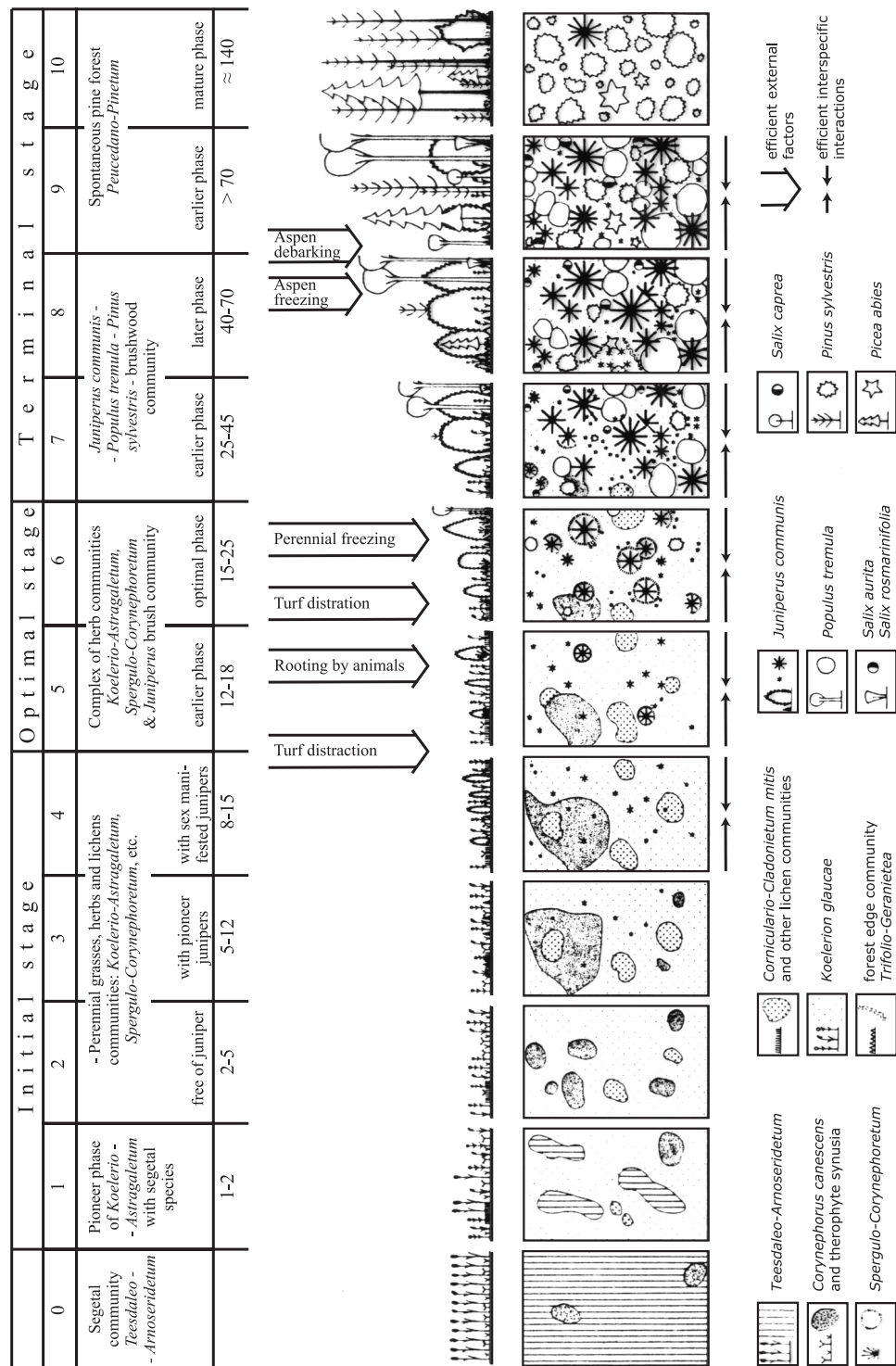


Fig. 2. Stages of secondary succession on the abandoned fields in Jelonka study area (based on FALINSKI *et al.* 1993).

and, similarly to the species known from the previous phase, they will be present till the phase of juniper-aspen brushwood (7-8). The proportion of psammophilous bryophytes, as well as lichens (mainly *Cladonia* spp.) increases. The mosaic of vegetation patches representing subatlantic pioneer psammophilous grassland with domination of therophytes (*Spergulo-Corynephorum* R. Tx. ex K. Czyżewska subass. *typicum*) or lichens (*S.-C.* subass. *cladinetosum mitis*) and with considerable participation of bryophytes (*S.-C. cladinetosum* var. with *Polytrichum piliferum*) (CZYŻEWSKA 1992), as well as patches of subcontinental psammophilous grassland of a *Koelerion glaucae* type is formed. Seedlings of juniper, pine and sometimes of aspen appear in greater quantities. At the end of the phase 4, which closes the initial stage of psammophilous grassland succession, forest mosses start to occur, e.g., *Sciuro-hypnum oedipodium*, *Hypnum cupressiforme* and *Dicranum scoparium*.

**Phases 5 and 6. A psammophilous herb-grassland and lichen communities in a complex including juniper shrub.** Duration 12-25 years after the field abandonment. An increase in the number of juniper as well as the growth in height and breadth of the juniper shrubs result in breakup of the lichen-moss sward and the formation of edge-strip aggregations of psammophilous therophytes around, comprising, e.g., *Teesdalea nudicaulis*, *Cerastium semidecandrum*, *Myosotis stricta*, *Spergula morisonii* as well as *Corynephorus canescens*. An increase in the projection cover of junipers and pines causes permanent shading of the ground, accumulation of needles and xeric relations in the soil, because of an almost complete interception of precipitation. A decrease in the number of species and frequency of mosses can be seen, as well as an intensive development of diverse lichen mats. In phase 6, several perennial species which attain their optimum in the brushwood phases, already occur, like *Peucedanum oreoselinum* and *Chamaecytisus ruthenicus*. Typical forest mosses appear around growing junipers and pine, aspen and spruce trees: *Dicranum polysetum* and *Hylocomium splendens*.

**Phase 7 and 8. A juniper-aspen brushwood with pine.** Duration 25-70 years after the field abandonment. Parallel processes of forming large juniper shrub aggregations, tree canopy closure, gradual disappearance of psammophilous communities and therophytes, as well as arrival of forest-edge perennials and willows take place. The brushwood becomes very dense and vertically complex, light and humidity conditions change, and there is an intensification of litter accumulation. The aspen and pine, accompanied by solitary birches and spruce, begin to form the tree stand. In the field layer, grassland species disappear and typical components of coniferous forests begin to occur, like *Melampyrum pratense*, *Vaccinium myrtillus*, *V. vitis-idaea*, *Pleurozium schreberi*, *Dicranum polysetum*, and *Hylocomium splendens*. Similarly, lichens typical of open vegetation are replaced by forest species. Temporal presence of, e.g., *Climacium dendroides*, *Brachythecium rutabulum*, *Plagiomnium affine*, the higro-hydrophilous bryophytes which have an ecological optimum in humid and wet sites, is characteristic of these phases. The juniper-aspen brushwood in the age of 60-70 years already attains stratification similar to that of a forest community in some places.

**Phase 9 and 10. Early and mature phase of stable pine forest.** Duration over 70 and about 140 years after the field abandonment, respectively. It is a subcontinental, mesophilous forest of *Peucedano-Pinetum* Mat. (1962) 1973 type with some subboreal components. The closest station of such forest, representing undisturbed final community of the chronosequence described from the Jelonka reserve, is situated ca 8 km to the northeast, in the Czechy Orlańskie reserve.

## 2. SUBJECT AND AIMS OF STUDY

Conversion of vast areas of natural forests into agriculture and subsequent abandonment resulting from a loss of productivity or socioeconomic and demographic reasons is one of the main causes leading to the decrease of biodiversity worldwide (DOBSON *et al.* 1997; STOATE *et al.* 2009). According to the afore mentioned authors, the European Union predictions are that up to 2030, land taken out of agricultural production will be one of the most striking features of the European landscape. The management of abandoned fields is one of the major problems which has to be urgently resolved in many countries in Europe, America and Asia (SMYKAŁA 1990; SOBCZAK 1990; SZUJECKI 1990; JOHNSTON & CROSSLEY 2002; DU *et al.* 2007; STOATE *et al.* 2009).

The main management policy to cope with abandoned farmland, especially on poor soils, is afforestation; this has positive impact for water, soil and air protection, enhances carbon sequestration thereby reducing atmospheric concentrations of carbon dioxide and leads to the increase of overall diversity (STOATE *et al.* 2009). However, it frequently meets serious difficulties and gives unsatisfactory results. The establishment and performance of outplanted seedlings has often been reported to depend on ectomycorrhizal fungi (PERRY *et al.* 1987; KROPP & LANGLOIS 1990; GROVE & LE TACON 1993; MARX *et al.* 2002; DUNABEITIA *et al.* 2004; RINCÓN *et al.* 2007), which may enhance uptake of water and nutrients (SMITH & READ 2008) and protect them against drought, pathogens and heavy metal pollution (MORIN *et al.* 1999; VAN TRICHELEN *et al.* 2001; ORTEGA *et al.* 2004). The soil cultivated for tens or even hundreds of years usually lacks a natural ECM inoculum (JOHNSON *et al.* 1991), thus the tree seedlings, either planted or regenerating spontaneously, may not find proper symbiotic partners. Without mycorrhizae on their roots, pine seedlings cannot survive on poor post-arable soil for more than 2 years (DOMINIK 1961).

It is well known that, apart from abiotic environmental conditions and the host plant presence, one of the most important causal factors affecting the onset and the subsequent process of fungal succession is the availability of species, either in form of spores, sclerotia, or mycelia (BRUNDRETT 1991; FRANKLAND 1992; TRAPPE & LUOMA 1992; JUMPPONEN *et al.* 2002; ASHKANNEJHAD & HORTON 2006; NARA 2006). This applies both to primary and secondary successional sites. In the absence of ECM inoculum, the development or recovery of ECM-dependent vegetation is possible only after the arrival of air-borne or animal-transported propagules from the 'outside', which depends on the inoculum source distance (site size) (DICKIE & REICH 2005) and time since disturbance (HELM *et al.* 1999; JUMPPONEN *et al.* 2002; JUMPPONEN 2003).

There is growing evidence from field studies, both in experimental and natural systems, that mycorrhizal symbionts can play a crucial role in successions of vegetation and vice versa (SMITH & READ 2008). Although there are a few studies of chronosequences of fungi in post-agricultural land developing after cultivation cessation (first generation or first rotation forests), almost all of them concern managed plantations of trees (MASON *et al.* 1982); some encounter such stands among others of different origin, e.g., together with primary stands or 2- and 3-rotation stands (TERMORSHUIZEN 1991). None of these studies include non-ECM fungi.

Fungal communities, both ECM and saprobic, associated with spontaneous and entirely natural and man-unaffected return of the forest onto the old-fields, forming a hundred years long chronosequence, have been examined in the present study for the first time.



The subject of the study is the succession of macrofungi on abandoned farmland and its relationship with the process of spontaneous, secondary succession of vegetation liberated from anthropogenic impact, leading to the restoration of the forest. The research comprised the observations of fungal carpophores as well as quantitative examination of ectomycorrhizal roots. The term *macrofungi* or *macromycetes* is used in accordance with ARNOLDS (1981, 1992a) and means all the fungi producing sporocarps visible by naked eye (usually larger than 1 mm).

It has been assumed that fungi, similarly to vascular plants, mosses and lichens, are permanent participants of the succession process studied. The aim of the research is to reveal the temporal patterns of macrofungal dynamics in abandoned fields and to relate them to the vegetation development. Thus, qualitative and quantitative examination of macromycetes in the range of successional phases of the series psammophilous grassland to mature pine forest was carried out, as well as basic quantitative analysis of ectomycorrhizal roots. Further the dynamics of fungal communities were related to changes in the composition and structure of phytocoenoses and habitat features; this allowed the addition of macrofungi to the model of old-field succession based on vegetation changes and lichen biota described above.

### 3. STUDY AREA

The research was carried out in two nature reserves, Jelonka and Czechy Orlańskie, situated near the southwest peripheries of the Białowieża old-growth forest (Fig. 1) in the complex of abandoned arable fields undergoing spontaneous secondary succession (Fig. 2).

**Geographical location, geological substrate and land surface features.** The study area is situated in mid-south part of the Równina Bielska plain (Nizina Północnopodlaska lowland, NE Poland), on the Precambrian East European platform, in the area of ground moraine plateau of the Middle Poland (Riss) glaciation. The most characteristic surface features are postglacial basins, hills of terminal moraine usually not exceeding 10 m, and a lot of big sandrs situated along the moraine, with some distinct cones usually over 180 m a.s.l. high (MOJSKI 1972). The Pleistocene deposits are mainly sands, gravels and glacial boulders. Among the Holocene deposits are sands forming indistinct ridges and organic sediments in the basins floor (FALIŃSKI *et al.* 1993).

**Climate.** The climate is determined by the situation of the area in NE Poland where the influence of the continental air is usually stronger than the influence of the maritime one. This results in big annual temperature amplitudes (22-23°C), long summers (over 90-100 days) and long cold winters (100-110 days). The vegetation season lasts ca 190 days. For about 140 days the minimal temperature drops below 0°C (ground frosts) (MARTYN 1995). The autumn ground frosts frequently repeating for several days cease vegetation many days prior to the average daily temperature attaining the 5°C threshold. The snow cover lasts over 90 days (PIÓRO 1973). The mean annual temperature is about 7°C. The temperature extremes registered in Białowieża (1949-2001) are +34.5°C and -38.7°C (FALIŃSKI 2002). The mean annual precipitation is 550-600 mm; the spring rainfall is usually more

abundant than the autumn one, which is also a feature of the continental climate (MARTYN 1995).

**Jelonka reserve.** The Jelonka reserve (N52.595; E23.364; alt. 178-185 m) is situated in the Kleszczów commune on either sides of the Hajnówka-Kleszczule road, about 1.2 km from the Jelonka village (see Fig.1). It remains under the administration of the Bielsk Podlaski Forest Inspectorate and comprises a compact area of 227 ha.

The reserve is situated in the catchments area of the Nurzec river (a tributary of the Bug river), but lacks any larger water courses. Two water-head areas – Trubianka Przednia and Trubianka Tylna lie to the south of the reserve. Within the reserve borders, three natural basins of probably ground-ice thawing origin are situated: Duży Ług (12 ha), Średni Ług (1.5 ha) and Mały Ług (0.3 ha). They form astatic water reservoirs with flat floors, filled with transitional peat bogs. They are part of the system of numerous drainageless basins occurring near the upper part of the Nurzec river (FALIŃSKI *et al.* 1993).

The reserve principally comprises former arable fields on poor mineral soils formed mainly from loamy sands. They are occupied with a complex of psammophilous grasslands, juniper shrubs, pine thickets and brushwood of juniper, aspen and pine (FALIŃSKI 1986b, 1998; FALIŃSKI *et al.* 1993). The occurrence and spatial distribution of these communities reflect the old ownership relations and time after abandonment. The peculiar landscape is made of narrow field-beds (5-20 m broad and 1-2 km long) set aside at different times beginning from World War I, representing different stages of secondary succession. However, as the habitat potential and environmental conditions show great homogeneity, they belong to the same successional series leading to the restoration of fresh pine forest *Peucedano-Pinetum*. In 1992, an area of about 70 ha was burned by an accidental fire and is now spontaneously regenerating.

The Jelonka reserve was established in 1989 for the strict protection of the vegetation complex formed on poor post-arable fields, which creates this unique landscape reflecting an old and traditional land use. The conservation extends throughout the whole ecological process shaping this landscape – the spontaneous secondary succession leading to natural forest return after centuries of deforestation and agricultural management (FALIŃSKI *et al.* 1993).

The area has been used for scientific research and education purposes mainly by the Białowieża Geobotanical Station of the Warsaw University (BSG UW). The monitoring of the succession process and studies on the ecology of pioneer species have been permanently carried out since 1971 (FALIŃSKI 1986a, b, 1998, 2002; set of plots no. 66) resulting, among other things, in the Faliński's model of the old-field secondary succession based on dynamics of vascular plants, supplemented with bryophytes and lichens by Czyżewska & Cieśliński (FALIŃSKI *et al.* 1993). After the fire in 1992, research on the spontaneous regeneration has also been undertaken (FALIŃSKI 1996, 2002; set of plots no. 79). Participation of macrofungi in this process was also examined (KAŁUCKA & SUMOROK 1996; SUMOROK 2001).

In the area of the present reserve, preliminary research on fungi accompanying *Juniperus communis* was performed by J.J. Barkman; after his untimely death a provisional list of the collected fungal species was published (BARKMAN & DE VRIES 1993).

**Czechy Orlańskie reserve.** The Czechy Orlańskie forest reserve (N52.631; E23.478; alt. 175 m) is situated 8 km to the northeast from the Jelonka reserve, in the neighbourhood of the Białowieża old-growth forest (see Fig. 1). It belongs to the Dubicze Cerkiewne

commune and is under the administration of the Bielsk Podlaski Forest Inspectorate. It occupies an area of 77.95 ha.

The reserve is situated on the flat plain between the Orlanka and Orla rivers in the catchments area of the river Narew.

It comprises a well preserved, compact fragment of pine and pine-spruce forests with a 100-year-old stand growing on post-arable land (FALIŃSKI 1992) probably as a result of semi-natural secondary succession (stimulated by leaving pine branches with cones on fresh fallow; Forestry Service, pers. comm., 1996). The following forest types can be distinguished:

- subcontinental mesophilous (fresh) pine forest *Peucedano-Pinetum* Mat. (1962) 1973 with subboreal features (the presence of *Goodyera repens* and a considerable amount of *Ptilium crista-castrensis*). A high proportion of *Chamaecytisus ratisbonensis* and *Lembotropis nigricans* provides evidence for the influence of subatlantic climate. Mainly the association of *P.-P. typicum* is represented but some patches of drier *P.-P. pulsatilletosum* can also be found;
- wet pine forest of *Molinio-Pinetum* Mat. 1973 type, which occupies small flat depressions. Oak trees as well as *Molinia caerulea* and *Vaccinium uliginosum* are present while spruce is almost lacking;
- spruce-pine forest representing the class and alliance of coniferous communities (*Vaccinio-Piceetea*, *Vaccinio-Piceetalia*), identified by FALIŃSKI (1992) as the *Quercus-Piceetum* (Mat. 1952) Mat. et Pol. 1955. These are rather transitional phytocoenoses as oak is completely lacking and the species typical of lowland spruce forests, like *Lycopodium annotinum*, *Goodyera repens*, *Moneses uniflora*, and *Plagiochila asplenioides* are very sparse.

In the area of the present reserve the stands were permanently but not intensively managed and harvested at places but never clear-felled (Plan Urządzenia Gospodarstwa... no date).

The reserve was established in 1995 for the conservation of the forests growing in between the Białowieża old-growth forest and historical Bielsk old-growth forest, as well as for permanent documentation of the processes shaping the forest communities in transitional phytogeographical location (FALIŃSKI 1992).

The Czechy Orłańskie forest reserve is an important element of the successional changes in the old-fields, documenting the final phase of the process of forest recovery (compare FALIŃSKI 1986a, b, 1998; FALIŃSKI *et al.* 1993). The BSG UW carried out long-term studies here using permanent plots (set of plots no. 128). In the 80ties of the last century, a series of experiments on accelerating and slowing down the successional processes were performed (FALIŃSKI 1992).

#### 4. MATERIAL AND METHODS

The research on the fungal succession was carried out during the years 1994-1998. The sporocarps of macromycetes were subject to qualitative and quantitative analysis in three consecutive seasons; the study focused on basidiomycetes as the proportion of ascomycetes fruit bodies was negligible. The ECM roots were also examined in two seasons. A series of soil analyses, as well as vegetation examination, were performed to document concurrent habitat features. Statistical analysis of the habitat-fungi relationships will be published separately.

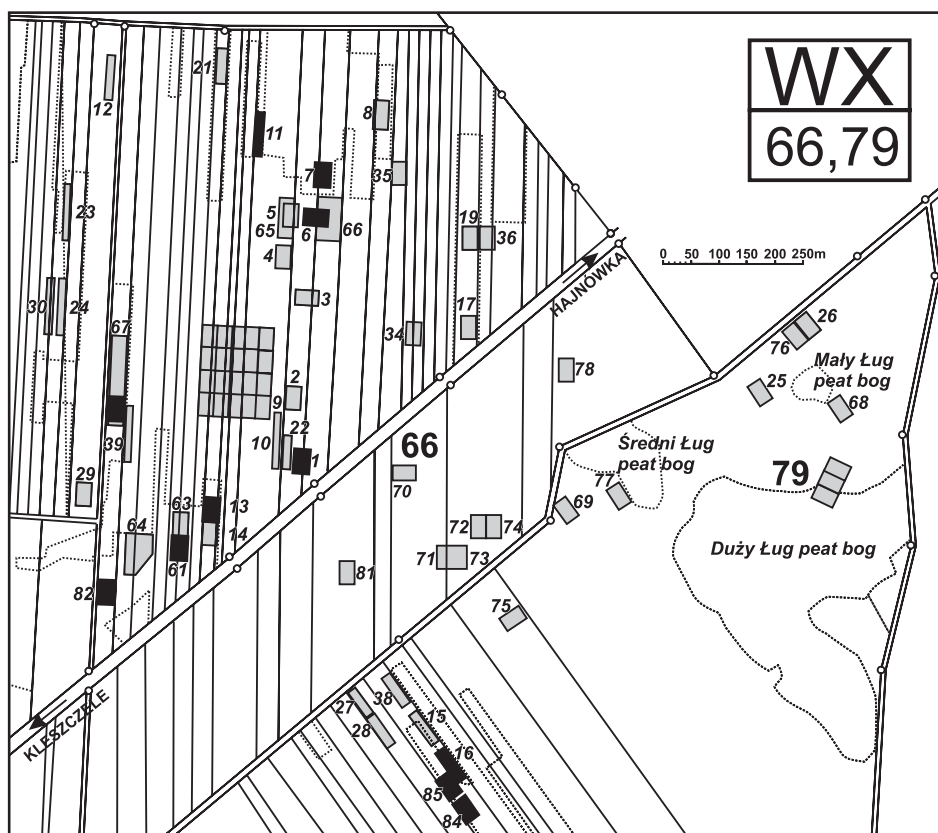


Fig. 3. Permanent plots within the Jelonka reserve and used in the studies of succession (based on FALIŃSKI 1998, augmented); the plots used in the present study are marked in black.

**Selection and distribution of observation plots.** The field observations were performed in permanent plots used by BSG UW for floristic-syndynamic research (FALIŃSKI 1977, 1980a, b, 1986a, b, 1998, 2002; FALIŃSKI *et al.* 1993). Most of them were set in the 1970s; some were established especially for the present research (nos 82, 84, 85). Each plot is marked with concrete posts at the corners, with BSG inscription and plot number, and is divided into subplots marked with wood stakes (see FALIŃSKI 1998, 2002).

The research was conducted in the Jelonka reserve, in 11 permanent plots, 1000 m<sup>2</sup> each (Fig. 3). They represent 6 phases of secondary succession (Fig. 4), and correspond to the initial, optimal and early terminal stage of vegetation development (according to the interpretation of aerial photographs by FALIŃSKI 1986a, b, reprinted in FALIŃSKI *et al.* 1993, and to the map by FALIŃSKI 1990 (1991), reprinted in FALIŃSKI 1998; updated by Faliński, pers. comm., 1994).

Five plots representing the phase of nearly mature pine forest of *Peucedano-Pinetum* type, situated in the Czechy Orlańskie reserve were also included, each of them of 400 m<sup>2</sup> (Figs 5, 6).

Each succession phase studied was represented by a total area of 2000 m<sup>2</sup>, with the exception of the earliest phase 2, for which only one 1000 m<sup>2</sup> plot could be found. Successional











Fig. 4. Permanent plots in the Jelonka reserve: plot 82 – phase 2 (A), plot 67 – phase 4 (B), plot 85 – phase 5 (C), plot 13 – phase 6 (D), plot 1 – phase 6/7 (E), plot 11 – phase 8 (F).

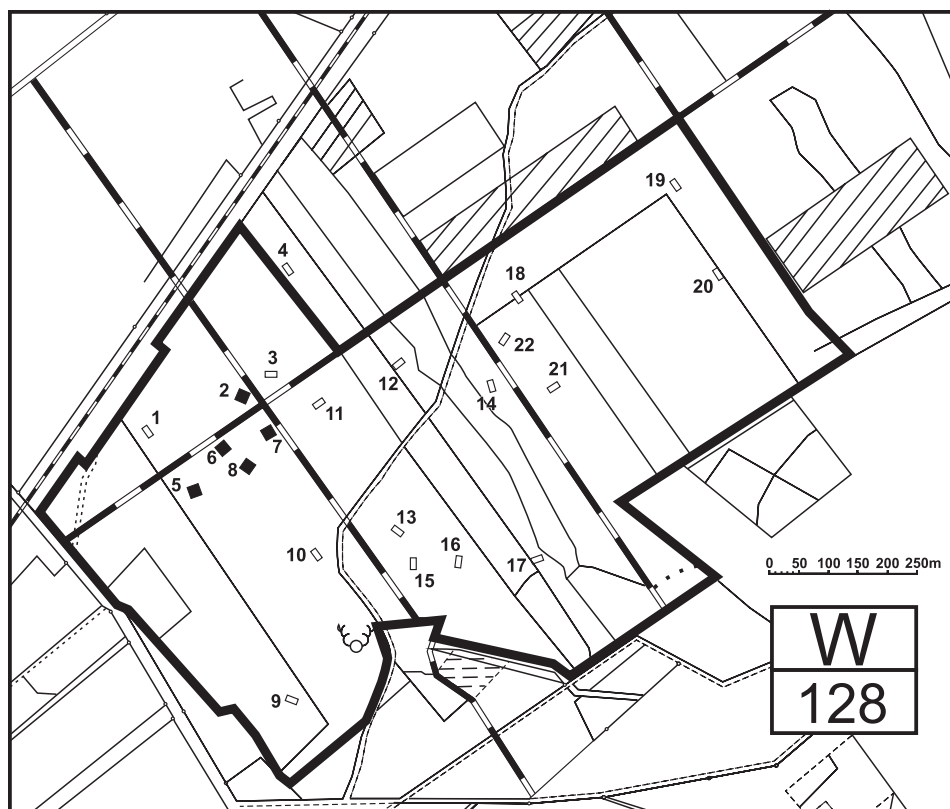


Fig. 5. Permanent plots in the Czechy Orlańskie reserve and used in the studies of succession (based on FALIŃSKI 1992, augmented); the plots used in the present study are marked in black.

status, size and shape, as well as internal division (subplots) of each plot are presented in Figure 7. In the present study, the original numbering of permanent plots is used.

**Soil description and analysis.** The examination of soil morphology was limited to the surface horizons as descriptions of full soil profiles were made in some plots of the Jelonka reserve in 1970s (FALIŃSKI 1987), as well as the analyses for forestry service purposes (Plan Urządzania Rezerwatu... no date). In all the profiles, the presence of plough humous horizon isolating deeper layers was clearly visible (Appendix 1).

In each succession phase, one plot was chosen for the description of soil morphology up to a depth of 50-60 cm. Samples for laboratory analyses were taken from each of the excavated horizons. The litter was sampled from the profile or from some place nearby, as in most plots it occurred patchily. The litter cover is included in the vegetation description (Appendix 2).

Additionally, ten samples of plough humous horizon (10-15 cm deep) were collected in each plot including a variety of sampling points differing in vegetation and litter. The samples originating from the same plot were mixed to obtain a bulk sample for the whole plot. In the pine forest of phase 9/10, ten samples were collected from all the 5 plots – two in each, and they were mixed together.



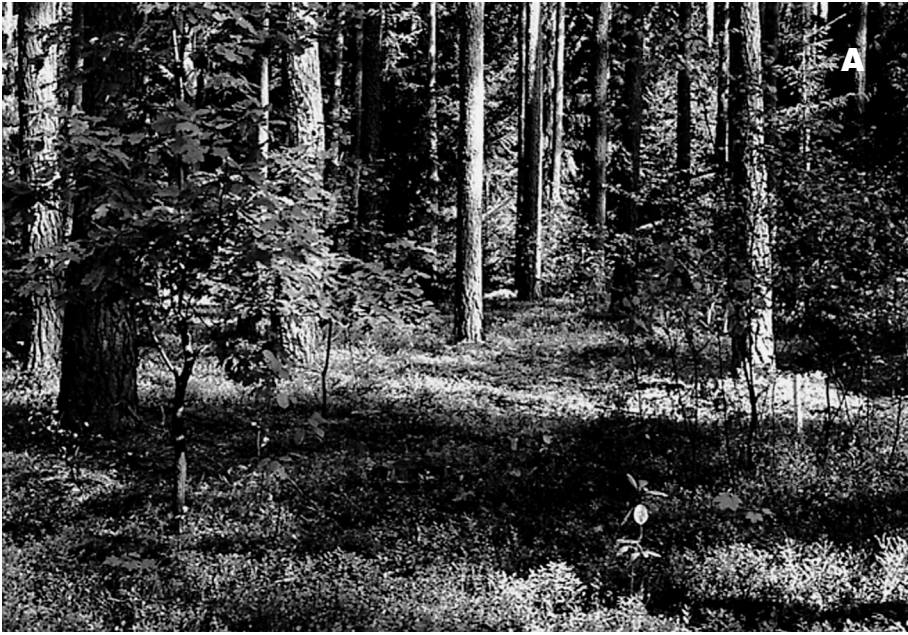


Fig. 6. Permanent plots in the Czechy Orlańskie reserve: plot 5 (A) and plot 6 (B), both representing phase 9/10.

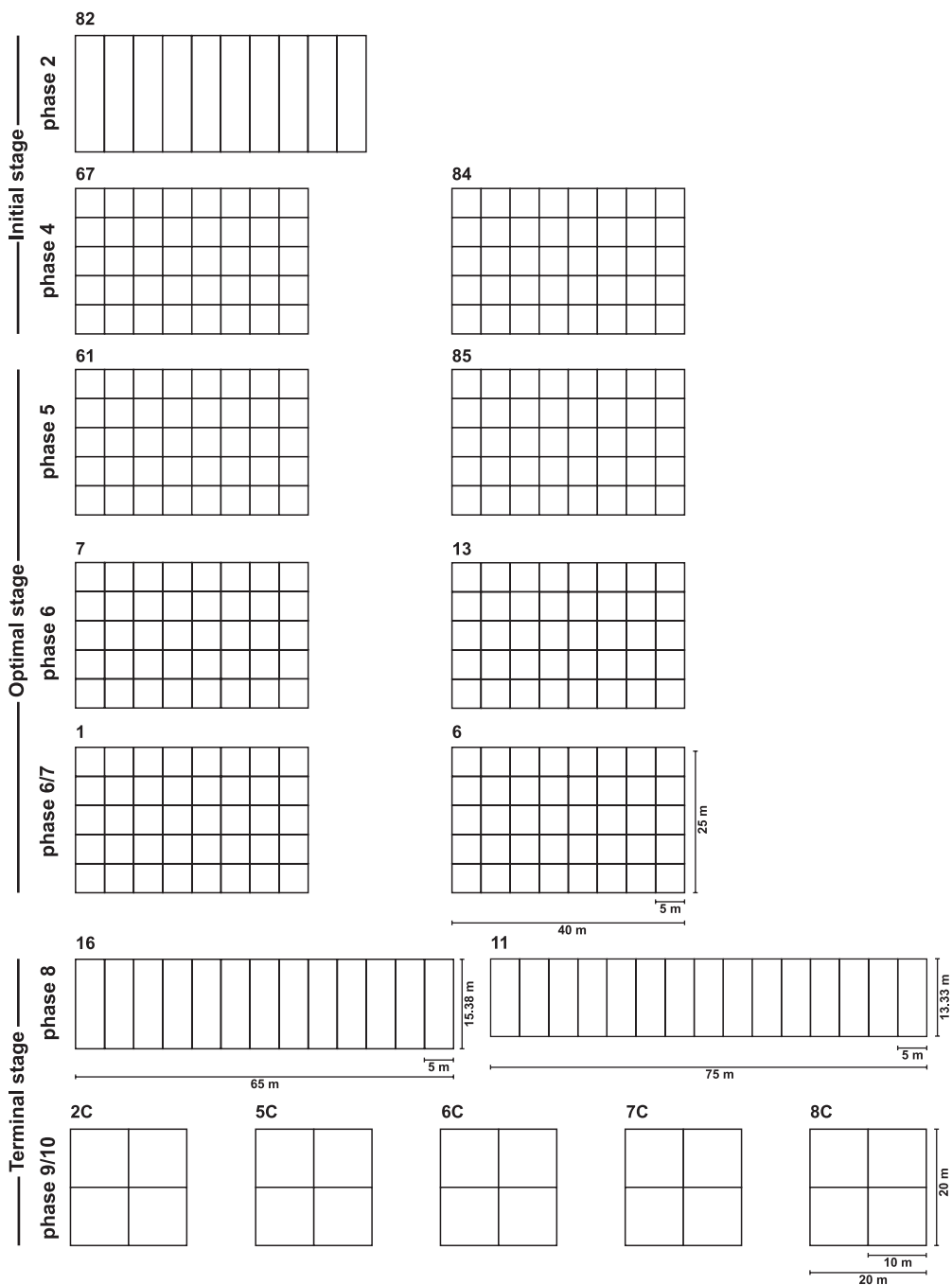


Fig. 7. Observation plots, showing numbering, internal division (subplots), and assignment to the succession phases and stages.

The soil analysis includes: (a) grain-size distribution in the samples from the soil profiles (according to the method by Cassagrande modified by Prószyński), (b) organic C content (lost-on-ignition method), (c) total N content (Kjeldahl's method), (d) pH in H<sub>2</sub>O and in KCl (1M).

Soil moisture was also analyzed in all the plots in 1997. Soil samples were collected in 5 terms (from July to November), at least 5 replicates in the 1000 m<sup>2</sup> plots and 2 replicates in 400 m<sup>2</sup> plots in each term.

**Vegetation description.** In the Jelonka reserve, the vegetation occupying the plots studied was described in June 1997, including the abundance of species according to the scale by LONDO (1975) (Appendix 2 – Tab. A2.1). The localization of relevés corresponds with the squares used for long-term phytosociological monitoring (see FALIŃSKI 1998, 2002).

The vegetation descriptions in the plots in the Czechy Orlańskie reserve are cited after FALIŃSKI (1992) (Appendix 2 – Tab. A2.2).

The nomenclature of vascular plants follows the work by MIREK *et al.* (2002), bryophytes – the work by OCHYRA *et al.* (2003), lichens – the work by SANTESSON *et al.* (2004).

Each plot was examined for the presence of coniferous and deciduous tree species. All the pines exceeding a height of 20 cm were counted and their age was estimated based on the number of branching points on the main stem. The age ranges of trees in plot descriptions refer to the third year of observations (1996).

**Mycological data – carpophores.** Systematic observations of fungal carpophores were carried out during three vegetative seasons (1994-1996), once a month on average, from May to November, depending on the weather conditions and intensity of sporocarp production. In the Jelonka reserve, each plot was visited 17 times, except for plot 82, which was established in 1995 and visited 13 times. In the Czechy Orlańskie reserve, 16 visits were performed for each plot. One by one, all the subplots in each plot were searched for macromycete carpophores during each visit; all the species were recorded and all the carpophores were counted.

*Qualitative analysis.* The carpophores collected were identified either fresh or dried according to the generally accepted macro- and microscopic methods using a NIKON SMZ-U binocular and NIKON E-400 light microscope.

The nomenclature of fungi follows KNUDSEN & VESTERHOLT (2008) for agaricoid and boletoid species, and LEGON *et al.* (2009) – for the other species. In case of species not included in the above mentioned sources, the nomenclature follows WOJEWODA (2003) and *Index Fungorum* (<http://www.indexfungorum.org>). The systematics of species is in accordance with KIRK *et al.* (2008).

All exsiccata have been deposited in the Herbarium Universitatis Lodzianensis (LOD).

In the field observations of fungal carpophores, taxonomical difficulties can cause problems with recording species and counting carpophores. In the present study, this was particularly true in case of the genera *Entoloma*, *Galerina* and *Cortinarius*. As these fungi occurred with high abundance and frequency, collective taxa were distinguished: *Entoloma* spp., *Galerina* spp. and *Cortinarius* spp. (see ARNOLDS 1992a, b). The latter genus was additionally divided into four groups of species based on the average dry weight of the carpophores:

<i>Cortinarius</i> spp. I	<0.2 g	<i>Cortinarius</i> spp. III	0.5-1.0 g
<i>Cortinarius</i> spp. II	0.2-0.5 g	<i>Cortinarius</i> spp. IV	>1.0 g

In the case of *Psilocybe montana*, part of the data refers to the rarer and less abundant species *P. phyllogena*; both species occurred in the same plots, their carpophores grew mixed together and clear distinction of the species in the field during dry weather conditions was impossible.

Forms and varieties were treated the same way as species in all quantitative analyses.

*Quantitative analyses.* The data on carpophore occurrence were analyzed using the following parameters (compare ARNOLDS 1981; KAŁUCKA 2008):

- abundance of carpophores of a species found in a plot during one observation season (year) (ACy);
- abundance of carpophores of a species found in a plot during the whole observation period (three years) (AC3y);
- average abundance of carpophores of a species in a plot per year (aACy);
- maximum abundance of carpophores per one observation (visit) (mACv); cf. maximum density of carpophores per visit (ARNOLDS 1981), maximum abundance = maximal number of carpophores per visit (KEIZER 1993a);
- spatial frequency (SF) – percentage of subplots where carpophores of a species were found; cf. spatial frequency (ARNOLDS 1981; JANSEN 1984);
- total temporal frequency (tTF) – number of observations in a plot when carpophores of a species were found; cf. number of occurrences (LISIEWSKA 1965, 1978; BUJAKIEWICZ 1981, 1991-1992), temporal frequency (ARNOLDS 1981);
- number of years when carpophores of a species were found (TFy).

In comparisons of fungal occurrence in the subsequent succession phases two parameters were applied (in the ECM and non-ECM species, separately):

- average annual density of carpophores of a species in the plots representing the same succession phase, per 1000 m<sup>2</sup> (aDCy) – the comparison shows the differences in carpophore production in the subsequent succession phases;
- average spatial frequency of a species in the plots representing the same succession phase (aSF) – the comparison shows the changes in spatial distribution of carpophores in the subsequent succession phases; as the number of plots as well as the number of subplots differ in the succession phases (Fig. 6), aSF were presented as the following classes: I = 1-10%, II = 11-20%, III = 21-30%, IV = 31-40%, V = 41-50%, VI = 51-60%, VII = 61-70%, VIII = 71-80%, IX = 81-90%, X = 91-100%.

For comparisons of species composition and diversity Sørensen's indices as well as Shannon's indices based on aDCy were calculated (KREBS 2001; ZAK & WILLIG 2004) for the ECM and non-ECM species, separately.

Species which occurred in only one succession phase (in one or more plots) were referred to as exclusive species for that phase.

Statistical analysis of the data on the occurrence of fungal species in the chronosequence studied was based on the total number of carpophores recorded in three seasons (DC3y) after the logarithmic transformation using the formula  $v = \ln(y + 1)$  (KEIZER 1993a) where  $v$  is the transformed value and  $y$  is the DC3y. The transformation reduces extreme values in the dataset and creates a scale similar to the one proposed by ARNOLDS (1981).

Groups (clusters) of species revealing similar patterns of abundance in particular periods of vegetation succession were distinguished with k-mean analysis (*STATISTICA* 9).

Mycological similarities of the plots and succession phases examined were analyzed with cluster analysis using the Euclidean distance as a similarity measure and Ward's method of grouping (*STATISTICA* 9).



*Analysis of the ECM fungi productivity.* The methods proposed by ARNOLDS (1981) were applied. For each species a specific, average value of the dry weight of a carpophore was assessed by weighing a number of representative, normally developed, carefully cleaned carpophores from the present study herbarial collection. According to ARNOLDS (1981), the results obtained using this method are only 3 to 5% higher than these obtained by weighing the carpophores dried at a temperature of 50°C. To calculate the average annual production of the ECM species, the specific dry weight of each species was multiplied by its average annual abundance.

It should be kept in mind that the production figures are underestimated; the number of observations and their frequency was not sufficient for recording completely all carpophores, some of them were omitted. Permanent or more frequent monitoring was not possible given the remoteness of the study area (450 km).

**Mycological data – ectomycorrhizal roots.** Quantitative examination of the ECM roots in the surface soil was carried out in September 1997 and in October 1998. In each plot in the Jelonka reserve (except for the plot 82 – phase 2) 6 soil cores were sampled (14.9 cm deep, 3.3 cm in diam; 127.44 cm<sup>3</sup>) in each season (12 samples all together). In the Czechy Orlańskie reserve, from the five 400 m<sup>2</sup> plots, 12 samples were collected. In the plot 82, where a single compact group of four young birches was growing (ca 1 m high), 3 soil cores were collected in each season at a distance of 0.5 m from the centre of the group.

The root samples were stored at 4°C before processing. They were washed carefully and all living roots of ECM trees (pine, aspen, willow, birch, spruce) were picked by hand under the dissecting microscope. The total length of fine roots (<2 mm; VOGT *et al.* 1983) was estimated using the line intercept method (TENNANT 1975; 2 mm grid). The mycorrhizal root tips were counted in two groups (KEIZER 1993a): (1) living tips with well developed mycorrhizal mantle, vividly coloured and firm, and (2) poor condition or dead tips with damaged or easily detachable mycorrhizal mantle, scruffy surface, usually dark brown colour. In the samples collected in 1998, mycorrhizal frequency was assessed (percentage of the number of living ECM tips in the total number of root tips). All the figures were calculated for 100 cm<sup>3</sup> of soil. Qualitative analysis of ectomycorrhizae was not within the scope of this project.

## 5. RESULTS

### 5.1. Fungi in permanent plots

#### Phase 2

**Plot 82.** Post-arable rusty soil; plough humous horizon (0-31 cm) of loamy sand; surface soil pH 5.5 (in H<sub>2</sub>O), C/N 14.4, median soil moisture in September-November varies between 9.67-18.15% (Appendix 1). Vegetation cover very dense (100% cover), consisting mainly of grass and herbs, with domination of *Poa pratensis*, *Agrostis capillaris*, *Stellaria graminea*, *Achillea millefolium*, *Vicia* spp., and with evenly scattered mosses (Appendix 2). Four solitary juniper specimens present (layer c/b). The plot is ectomycorrhizal (ECM)

tree-free with the exception of four birches growing in one group ca 1 m high. Although the cultivation practices in this plot have been ceased for more than 30 years (Forestry Service, pers. com., 1996), vegetation succession is still in its early initial stage. Elsewhere in the area studied, phase 2 usually lasts no longer than 2-5 years after the field abandonment (FALIŃSKI *et al.* 1993; Fig. 2).

In the plot representing phase 2 of the vegetation succession 15 basidiomycete species were found (Appendix 3, Tab. A3.1). All the species recorded were saprotrophic, growing on dead plant remnants or on soil. No ECM fungi were found despite the presence of a group of four young birches in the plot.

Dense field layer reaching the height of 0.5 m did not favour the growth of macromycete carpophores. Infrequent soil inhabiting species which emerged from among the compact turf of living and dead plant roots and shoots possessed rather firm and elastic carpophores (*Marasmius oreades*) or they occurred in places occupied by mosses (*Cystoderma amianthinum*, *Clitocybe diatreta* and *Mycena pura*). The majority of species inhabited plant remains at different stages of decay. They belong to the genera *Coprinopsis*, *Crinipellis*, *Entoloma*, *Mycena* and *Parasola*. *Coprinopsis friesii* and *Parasola plicatilis* are species exclusive to phase 2, attaining relatively high frequency there.

The feature characteristic of this phase is an ephemeral occurrence of fungi. Only two species – *Marasmius oreades* and *Crinipellis scabella* were producing sporocarps regularly (temporal frequency 10 and 6 in 13 observations, respectively); the other species occurred sporadically. Some of them were fruiting occasionally but then very abundantly, e.g., apart from *Marasmius oreades* and *Crinipellis scabella* – *Mycena olivaceomarginata*, *M. flavoalba* and *M. leptcephala*. Usually, a high abundance of carpophores was accompanied by a high frequency.

#### Phase 4

**Plot 67.** Post-arable rusty soil; plough humous horizon present; surface soil pH 4.6 (in H<sub>2</sub>O), C/N 16.1, median soil moisture in September-November varies between 6.88-14.04% (Appendix 1). Row litter cover of a mosaic character, on average ca 20%. Psammophilous grassland with high cover of moss layer and patches of lichens, with scattered juniper, birch, aspen and willow, and pine mainly in *b* layer (Appendix 2). Pines 116/1000 m<sup>2</sup>, present in 35/40 subplots (87.5%), 0-8 per subplot, most of 6-8 years old, the oldest specimens 11-12 years old. One spruce tree (1m high) present. Deciduous trees present in 32/40 subplots (80%). The plot is situated in the vicinity of juniper-aspen brushwood.

**Plot 84.** Post-arable rusty soil; plough humous horizon (0-28 cm) of loamy sand; surface soil pH 4.8 (in H<sub>2</sub>O), C/N 14.5, median soil moisture in September-November varies between 5.02-14.85% (Appendix 1). Row litter present only at places, under the oldest tree groups. Psammophilous grassland with high cover of moss layer, with considerable presence of juniper, birch, aspen, willow and pine mainly in *b* layer (Appendix 2). Pines 147/1000 m<sup>2</sup>, present in 38/40 subplots (95%), 0-11 per subplot, most 6-8 years old, the oldest specimens 12(15) years old. Three young spruce trees present in one subplot. Deciduous trees present in 40/40 subplots (100%). The plot is situated in close vicinity to juniper-aspen brushwood.

Ninety two species of macromycetes were found in the plots representing phase 4, among them 44 ECM species (47.8%), 47 saprotrophic species (51.1%), and 1 species of parasite (1.1%) (Appendix 3, Tab. A3.2, A3.3). Species similarity (based on Sørensen's

index) between the examined plots amounts to 64.6% in the ECM fungi and 67.6% in the other species. The respective similarity for the aggregated group of vascular plants, bryophytes and lichens amounts to 70.9%.

The majority of the ECM fungi are symbionts of pine. Among them *Inocybe lacera* is a dominating species; *Suillus luteus* and *Amanita muscaria* also occur with great abundance. Participation of *Laccaria proxima*, as well as of species of the genus *Cortinarius* producing tiny sporocarps, is distinctive. *Cortinarius cinnamomeoluteus* occurs regularly and with high abundance. The species mentioned fruit both with high abundance and frequency (over 50%).

Apart of species associated with pine, ECM symbionts of deciduous trees also occurred, e.g., *Leccinum aurantiacum* under aspen and *Lactarius vietus*, *Russula subfoetens* and *Leccinum scabrum* under birch. Although the spatial frequency of these trees in the plots representing phase 4 equals that of pine, and they are also present in the neighbouring juniper-aspen brushwood, the participation of the ECM fungi specific to these hosts is relatively low, both in terms of species numbers and carpophore abundance and frequency.

Among the saprotrophic fungi, species growing on soil, frequently among mosses, prevail, like *Psilocybe montana*, *Clitocybe diatreta* and species of *Entoloma*.

As litter accumulation in the examined plots is low, species growing on litter and wood occur with low frequency and abundance. There are some exceptions – the species growing on the remains of herbs and grasses, such as *Mycena olivaceomarginata*, *M. flavoalba* and *M. leptcephala*, as well as *Mycetinis scorodoni*, the only species attaining relatively high abundance and frequency exclusively within phase 4. Also, *Gymnopus androsaceus* inhabiting pine needles under some of the oldest trees (12-15 years old) was recorded as well as sparse carpophores of *Gymnopus dryophilus* on fallen leaves of deciduous trees and *Flammulaster granulatus* on fallen twigs. Among the sporadic species growing on wood were *Dacrymyces stillatus*, *Sphaerobolus stellatus*, *Tubaria confragosa* (on pine twigs), *Crepidotus variabilis* (on dead heather parts), *Hypholoma fasciculare* (on the wood stake marking the plot).

Relatively abundant were the carpophores of *Galerina* spp., inhabiting moss tufts, mainly *Polytrichum juniperinum* and *P. commune*. *Cantharellula umbonata*, the moss parasite was also found there.

## Phase 5

**Plot 61.** Post-arable rusty soil; plough humous horizon (0-24 cm) of loamy sand; surface soil pH 4.8 (in H<sub>2</sub>O), C/N 16.2, median soil moisture in September-November varies between 4.76-14.91% (Appendix 1). Litter cover of a mosaic character but considerable, on average ca 40%, consisting mainly of loose pine and juniper needles. Psammophilous grassland with abundant presence of mosses and high cover of lichens, almost devoid of aspen, with scattered birch, evenly distributed juniper and pine mainly in *b* layer (Appendix 2). Pines 105/1000 m<sup>2</sup>, present in 39/40 subplots (97.5%), 0-6 per subplot, most 8-10(12) years old, the oldest specimens 15-17 years old; numerous seedlings. Deciduous trees present only in 5/40 subplots (12.5%). The ground disturbed by wild boars at places.

**Plot 85.** Post-arable rusty soil; plough humous horizon present; surface soil pH 4.8 (in H<sub>2</sub>O), C/N 14.8, median soil moisture in September-November varies between 3.78-12.37% (Appendix 1). Row litter present only at places, under the oldest tree groups. Psammophilous grassland with considerable presence of mosses and lichens, with

dense and evenly distributed juniper shrubs, aspen, scattered birch, and pine mainly in *b* layer (Appendix 2). Pines 72/1000 m<sup>2</sup>, present in 28/40 subplots (70%), 0-4(10) per subplot, most 9-11 years old, the oldest specimens 12-16 years old. Deciduous trees present in 31/40 subplots (77.5%). The plot is situated in close vicinity to juniper-aspen brushwood.

A total number of 96 species of fungi were found in the plots representing phase 5 of the vegetation development, among them 39 ECM species (40.6%), 56 saprotrophic (58.3%) and 1 parasitic species (1.0%) (Appendix 3, Tab. A3.4, A3.5). Species similarity (based on Sørensen's index) between the examined plots amounts to 63.2% in the ECM fungi and 56.4% in the other species. The respective similarity in the aggregated group of vascular plants, bryophytes and lichens amounts to 58.2%.

Similarly to phase 4, in the ECM fungi the species typically associated with pine prevail. The most abundant is *Cortinarius cinnamomeoluteus*. Numerous carpophores are still produced by *Inocybe lacera* although its participation, both in terms of abundance and frequency, decreases. Among dominating species are also *Laccaria proxima* and *Hygrophorus hypothejus*; high abundance and the highest frequency is attained by *Suillus luteus*.

A considerable share of deciduous trees in plot 85 and in its close vicinity results in the presence of *Leccinum aurantiacum*, *Lactarius uvidus*, *Russula clavipes*, *R. galochroa* and *R. subfoetens* – species specific to aspen and birch, however, they produce only single carpophores close to the bordering juniper-aspen brushwood.

In saprotrophic fungi the same species as in phase 4 dominate – *Clitocybe diatreta* and *Psilocybe montana*. Most of the species fruit on soil. The occurrence of a number of gasteromycetoid species, mainly *Lycoperdon*, is specific for this phase. The species inhabiting litter and wood are rather sporadic, only *Mycena leptcephala* and *M. flavoalba* produce abundant carpophores on decaying plant remnants and *Collybia cirrhata* on old sporocarps of other fungi.

Lower cover of moss layer in the plots of phase 5 comparing to the plots of the phase 4 results in a decreased abundance of *Galerina* species and of *Cystoderma amianthinum* and *Cantharellula umbonata*.

## Phase 6

**Plot 7.** Post-arable rusty soil; plough humous horizon present; surface soil pH 4.7 (in H<sub>2</sub>O), C/N 17.2, median soil moisture in September-November varies between 5.73-13.53% (Appendix 1). Litter cover uneven, almost lacking at places but in many subplots exceeding 70%, consisting mainly of loose pine and juniper needles and twigs; in the SE part of the plot considerable amounts of aspen leaves from the neighbouring aspen brushwood. Psammophilous grassland with considerable moss cover, with scattered solitary birch and aspen, even and dense juniper shrubs and pine both in *a* and *b* layer (Appendix 2). Pines 131/1000 m<sup>2</sup>, present in 40/40 subplots (100%), 1-8 per subplot, most 8-15 years old, the oldest specimens 17-19 years old. Deciduous trees present only in 6/40 subplots (15%). The plot is situated in close vicinity to juniper-aspen brushwood.

**Plot 13.** Post-arable rusty soil; plough humous horizon (0-25 cm) of loamy sand; surface soil pH 4.8 (in H<sub>2</sub>O), C/N 18.4, median soil moisture in September-November varies between 6.00-10.64% (Appendix 1). Litter cover exceeding 70% in most of the plot, consisting mainly of loose pine and juniper needles and twigs, also with heather remnants at places. Psammophilous grassland with moderate moss and lichen cover, with a few solitary



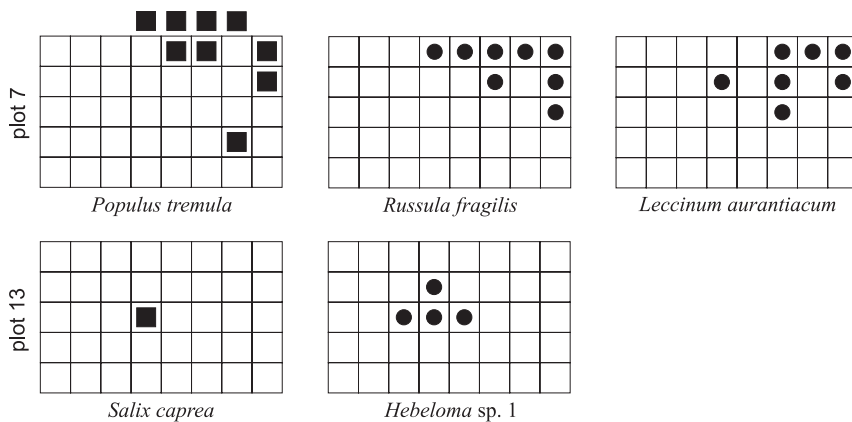


Fig. 8. Distribution of *Populus tremula* and *Salix caprea* and associated ECM fungi in the plots of phase 6.

aspens, two oak seedlings and one *Salix caprea*, even and dense juniper shrubs, and pine both in *a* and *b* layer (Appendix 2). Pines very dense, 227/1000 m<sup>2</sup>, present in 38/40 subplots (95%), 0-15(21) per subplot, most 8-13(15) years old, the oldest specimens 17-19 years old; numerous seedlings. Deciduous trees present only in 6/40 subplots (15%). The ground disturbed by wild boars at places.

A total number of 114 species of fungi were found in the plots representing phase 6 of the vegetation development, among them 47 ECM species (41.2%), 66 saprotrophic (57.9%) and 1 parasitic species (0.9%) (Appendix 3, Tab. A3.6, A3.7). Species similarity (based on Sørensen's index) between the examined plots amounts to 76.3% in the ECM fungi and 63.3% in the other species. The respective similarity in the aggregated group of vascular plants, bryophytes and lichens amounts 66.1%.

The spatial frequency of deciduous trees does not exceed 15% in both of the plots. The carpophores of species like *Leccinum aurantiacum*, *Russula fragilis*, *Hebeloma sp. 1* were found only in close vicinity of their probable host-trees (Fig. 8).

Mycorrhizal symbionts of pine are dominating, among them *Hygrophorus hypothecus*, *Cortinarius cinnamomeoluteus* and other species of *Cortinarius* producing small carpophores, *Amanita muscaria*, *Suillus luteus*, *Chalciporus piperatus*, *Laccaria proxima*, *Tricholoma albobrunneum* and *T. portentosum*. The species mentioned attain very high abundance and frequency. Abundant occurrence is also a feature of *Thelephora terrestris*, *Inocybe lacera*, *Lactarius rufus*, *Tricholoma equestre*, *Cortinarius parevernus* and *Laccaria bicolor*. It is striking that in both plots of phase 6 a number of species of *Tricholoma* raise their spatial frequency and start to produce conspicuously more carpophores.

The ECM species exclusive for this phase are very few and occur sporadically.

Dominant saprotrophic species are still, as in the previous phase, *Clitocybe diatreta*, *Psilocybe montana*, *Cystoderma amiantinum* and *Rickenella fibula*. Considerable bryophyte cover and a large proportion of mosses forming high tufts (*Polytrichum juniperinum*, *P. commune*) result in high abundance and frequency of species of *Galerina*. There is an increase in species richness and in carpophore abundance of species growing on litter, such as *Gymnopus androsaceus* and *Ampulloclitocybe clavipes*. The fruit bodies of species of *Clitocybe* and *Mycena* were also found more frequently on litter than on soil. For the first

time in the chronosequence studied, there occur *Strobilurus stephanocystis* and *Baeospora myosura*, the species inhabiting pine cones.

An increasing amount of dead wood, mainly of fine woody debris, results in growing diversity of fungi on this kind of substrate – on fallen twigs, partly buried wood pieces, on dead lower branches of trees. Among them are *Dacrymyces variisporus*, *Flammulaster granulosus*, *Trichaptum fuscoviolaceum*, *Leptosporomyces galzinii*, *Calocera furcata*, *Sphaerobolus stellatus*, *Pholiota gummosa*, *Tremella mesenterica*, *Phanerochaete sanguinea*, *Hypholoma fasciculare*.

#### Phase 6/7

**Plot 1.** Post-arable rusty soil; plough humous horizon (0-30 cm) of loamy sand; surface soil pH 4.8 (in H<sub>2</sub>O), C/N 17.7, median soil moisture in September-November varies between 4.24-10.20% (Appendix 1). Litter cover exceeding 70% in most of the plot, tending to accumulate in depressions, consisting almost exclusively of loose pine and juniper needles, twigs and branches, also with heather remnants at places. Patches of psammophilous grassland with moderate moss and lichen cover, with a few sparse aspens, even and dense juniper shrubs, and pine thickets both in *a* and *b* layer (Appendix 2). Pines 151/1000 m<sup>2</sup>, present in 39/40 subplots (97.5%), 0-9 per subplot, most 12-15(17) years old, the oldest specimens 22-24 years old; numerous seedlings. Solitary aspens present only in 2/40 subplots (5%). The ground disturbed by wild boars at places.

**Plot 6.** Post-arable rusty soil; plough humous horizon (0-20 cm) of loamy sand; surface soil pH 4.8 (in H<sub>2</sub>O), C/N 15.7, median soil moisture in September-November varies between 4.00-15.18% (Appendix 1). Thick litter cover exceeding 80% in most of the plot, tending to accumulate in depressions, consisting almost exclusively of loose pine and juniper needles, twigs and branches, also with heather remnants at places. Patches of psammophilous grassland with moderate moss and lichen cover, even and dense juniper shrubs, and pine thickets both in *a* and *b* layer (Appendix 2). Pines very dense, 257/1000 m<sup>2</sup>, present in 37/40 subplots (92.5%), 0-11(21) per subplot, most 8-15(18) years old, the oldest specimens 23-25 years old; numerous seedlings. No deciduous trees present.

A total number of 105 species of fungi were found in the plots representing phase 6/7 of the vegetation development, among them 42 ECM species (40.0%) and 63 saprotrophic species (60.0%) (Appendix 3, Tab. A3.8, A3.9). Species similarity (based on Sørensen's index) between the examined plots amounts to 83.3% in the ECM fungi and 66.0% in the other species. The respective similarity in the aggregated group of vascular plants, bryophytes and lichens amounts to 73.6%.

Nearly all the ECM fungi recorded in the examined plots are associated with pine which is a dominant tree in stand composition. The most abundant are *Cortinarius cinnamomeoluteus* and other species of *Cortinarius* producing small fruit bodies, *Hygrophorus hypothejus*, *Laccaria proxima*, *Amanita muscaria*, *Lactarius rufus*, *Tricholoma portentosum*, *T. equestre* and *T. albobrunneum*. The carpophores of *Suillus luteus* and *Tricholoma saponaceum* var. *saponaceum* are very numerous; relatively frequent are also *Boletus edulis*, *Cortinarius anthracinus* and *Sarcodon squamosus*. However, the abundance of many ECM fungi decreases.

A particularly high species richness as well as abundance and frequency of the genus *Tricholoma* is characteristic of this phase of vegetation development.

The species composition of saprotrophic fungi does not differ very much from the previous phase; *Clitocybe diatreta* and *Psilocybe montana* are dominating, among the abundant taxa are also *Gymnopus androsaceus* and species of *Galerina*, *Collybia cirrhata* and *Clitocybe vibecina*. A great variety of microhabitats favour the occurrence of many species growing on soil, litter, wood and among bryophytes. As in the ECM fungi, many saprobic species that occurred with high frequency and produced abundant carpophores in earlier phases seem to be less productive.

Exclusive species for phase 6/7 are almost completely lacking.

#### Phase 8

**Plot 11.** Post-arable rusty soil; plough humous horizon present; surface soil pH 5.3 (in H<sub>2</sub>O), C/N 21.0; thin humus accumulation horizon present, beginning of the podzolization process visible; median soil moisture in September-November varies between 7.12-17.89% (Appendix 1). Litter layer cover reaching almost 100%, consisting mainly of aspen leaves mixed with pine and juniper needles, rich in fallen twigs and branches, in different stages of decay. Dense brushwood of aspen, pine, birch and *Salix caprea* in *a* layer, with thick undergrowth of juniper and broadleaves, rather poor herb layer and scattered mosses (Appendix 2). Pines not numerous but evenly distributed, 34/1000 m<sup>2</sup>, present in 15/15 subplots (100%), 0-5 per subplot, most 20-30 years old, the oldest specimens more than 30-35 years old. One spruce tree present. Deciduous trees present in all subplots (100%).

**Plot 16.** Post-arable rusty soil; plough humous horizon (1.5-25 cm) of loamy sand; surface soil pH 5.0 (in H<sub>2</sub>O), C/N 17.8; thin humus accumulation horizon (0-1.5 cm) present, beginning of the podzolization process under coniferous trees visible; median soil moisture in September-November varies between 6.66-20.10% (Appendix 1). Litter layer cover reaching almost 100%, consisting mainly of aspen and birch leaves, juniper needles, rather sparse pine and spruce needles, and herbaceous plant debris, rich in fallen twigs and branches, in different stages of decay. Dense brushwood consisting mainly of aspen mixed with solitary birches, pines and spruces in *a* layer, with thick undergrowth of juniper and broadleaves (among them 3 willow species), high cover of herb layer and scattered mosses (Appendix 2). Pines rather few, 6/1000 m<sup>2</sup>, present in 6/13 subplots (46%), 0-1 per subplot, most 20-30 years old, the oldest specimens more than 30-35 years old. Three spruce trees present. Deciduous trees present in all subplots (13/13, 100%).

A total number of 183 species of fungi were found in the plots representing phase 8 of the vegetation development, among them 49 ECM species (26.8%) and 134 saprotrophic species (73.2%) (Appendix 3, Tab. A3.10, A3.11). Species similarity (based on Sørensen's index) between the examined plots amounts to 60.9% in the ECM fungi and 64.6% in the other species. The respective similarity in the aggregated group of vascular plants, bryophytes and lichens amounts to 66.2%.

The juniper-aspen brushwood is a transitional vegetation community between psammophilous grassland complex and fresh pine forest. The tree stand in the plots representing phase 8 comprises a very high proportion of deciduous species, mainly aspen but also birch, willow and oak, and relatively scattered conifers. However, in the carpophore community of the ECM fungi both groups of symbionts are well represented – of deciduous and of coniferous trees. The former consists of, e.g., *Cortinarius trivialis*, *C. alboviolaceus*, *Russula galochroa*, *R. versicolor*, *Leccinum aurantiacum*, *L. scabrum*,

*Lactarius torminosus*, *L. uvidus*, *Inocybe flocculosa*, *Amanita phalloides*, and many species of *Cortinarius*. The latter is represented by, e.g., *Tricholoma portentosum*, *T. imbricatum*, *T. albobrunneum*, *Chalciporus piperatus*, *Suillus luteus*, *Cortinarius cinnamomeoluteus* and *Russula paludosa*. Many of the ECM species found are host-generalists, like *Amanita muscaria*, *A. pantherina*, *A. rubescens*, *Boletus edulis*, *Clitopilus prunulus*, *Paxillus involutus*.

The ECM fungal community of phase 8 comprises a numerous group of exclusive species, among them *Cortinarius trivialis* and *C. albobviolaceus*, a majority of the species from the *Cortinarius* spp. II and *Cortinarius* spp. III complexes (producing medium-sized and big carpophores), *Hebeloma* sp. 3, *Lactarius torminosus*, *Inocybe flocculosa* as well as *Amanita gemmata*, *A. phalloides*, *Hebeloma stenocystis*, *Inocybe* cf. *margaritispora*, *Lactarius necator*, *Russula* cf. *mustelina*, *R. versicolor*. The species listed first occur with high abundance and/or frequency, in contrast to the exclusive species of the earlier phases of succession. *Cortinarius trivialis*, *C. albobviolaceus* and *Lactarius torminosus* were found in both plots representing phase 8.

Twenty three ECM species which were present in the earlier phases of the succession produce their sporocarps in phase 8 but are not present in the subsequent phase – the pine forest. The optimum of occurrence of most of them in terms of abundance and frequency of carpophores coincides with the previous succession stage (the optimal stage). The optimum in phase 8 is attained only by *Leccinum aurantiacum*, *Clitopilus prunulus*, *Russula clavipes*, *R. galochroa* and *Lactarius uvidus*, these species occurring sporadically in the communities other than juniper-aspen brushwood and always in its close neighbourhood.

Three species – *Amanita citrina*, *Cantharellus cibarius* and *Russula paludosa* start fruiting in this phase for the first time and they continue their occurrence in the next phase – the pine forest.

Among the dominant ECM species producing large numbers of carpophores are *Cortinarius trivialis* and other species of small and medium-sized *Cortinarius*, *Amanita citrina* and *Cantharellus cibarius*.

The most abundant saprotrophic species are those which usually inhabit an accessible substrate (fallen dead branches, twigs, litter) in a plentiful way – *Dacrymyces stillatus*, *Gymnopus androsaceus*, *Crepidotus lundellii* and *Tubaria furfuracea*. Relatively abundant are also *Mycena pura*, *M. leptcephala*, *M. olivaceomarginata*, *Galerina hypnorum*, *Clitocybe candicans*. However, most of them occurred also in earlier phases and some even produced larger number of carpophores then.

The participation of saprobic species growing on soil is rather low – 17%, of species occurring among mosses – less than 3%; ca 80% are the litter-inhabiting and lignicolous fungi in equal proportion.

Nearly a half of the saprotrophic species found in the plots of phase 8 are exclusive taxa, some of them are relatively frequent in both of the plots, e.g., *Coprinellus xanthothrix*, *Flammulaster rhombosporus*, *Mycena stipata* and *Psathyrella candolleana*. This probably results from very specific microclimate conditions, the highest participation of deciduous trees and plentiful, highly diverse substrates.

The characteristic feature of species composition, both ECM and saprobic fungi, is also the presence of species typical of rather humid or even wet and boggy places, like *Russula paludosa*, *Marasmius epiphyllus*, *Gymnopus aquosus*, *Mycena supina*, *Phaeomarasmium erinaceus* and *Lentinus suavisissimus*.

**Plots 2C-8C.** Podzolic soil developing on post-arable rusty soil, plough humous horizon (20-35 cm) still present. Surface soil pH 4.6 (in H<sub>2</sub>O), C/N 24.0, median soil moisture in September-November varies between 11.05-24.54%. Litter layer on average 6 cm thick, differentiated (Ol, Of, Oh), consisting mainly of pine needles, bark, numerous fallen twigs and branches (Appendix 1). Nearly mature fresh pine forest, with scattered spruce, birch, oak and willow trees, poor undergrowth with dying junipers, rich field layer with prevailing dense shrubs of perennials and nearly 100% cover of mosses (Appendix 2). Pines mainly even-aged, over a hundred years old, on average 40/1000 m<sup>2</sup>, evenly distributed. Numerous pine seedlings. The ground disturbed by wild boars at places.

A total number of 98 species of fungi were found in the plots representing the phase 9/10 of the vegetation development, among them 30 ECM species (30.6%), 67 saprotrophic species (68.4%) and 1 species of parasite (Appendix 3, Tab. A3.12-16). Species similarity (based on Sørensen's index) between the examined plots amounts to 65.0-78.9% in the ECM fungi and 54.2-65.6% in the other species. The respective similarity in the aggregated group of vascular plants, bryophytes and lichens amounts to 71.7-81.6%.

The species composition of the ECM fungi is typical of fresh pine forest. Among dominating species are *Paxillus involutus*, *Amanita citrina*, *Cortinarius semisanguineus* and *C. caperatus*. Also, *Russula emetica*, *Lactarius rufus* and *Thelephora terrestris* occur with high abundance.

About half of the ECM species were recorded in the earlier succession phases and most of them were also more abundant there, e.g., *Amanita citrina*, *Lactarius rufus*, *Thelephora terrestris*, *Cantharellus cibarius*, *Laccaria proxima*, *Russula fragilis* and *Amanita rubescens*. However, the abundance of carpophores and spatial frequency of *Paxillus involutus*, *Russula emetica*, *R. paludosa* and *Xerocomus badius*, is markedly higher in the forest plots of phase 9/10.

The other half of the ECM species are exclusive species for that phase; these are typical forest species growing under conifers, like *Cortinarius caperatus* and *Amanita porphyria*, as well as many *Russula* species, the genus best represented here.

The saprotrophic species grow mainly on pine litter and woody debris, and frequently occur among mosses. Among the species dominating in terms of abundance are *Dacrymyces stillatus*, *Gymnopus androsaceus*, *Collybia cirrhata* and *Mycena zephirus* – the fungi in which massive production of small carpophores is a biological feature of the species. In some species, like *Leptosporomyces galzinii*, *Mycena metata*, *M. galopus*, *Galerina pumila*, *Cystoderma amiantinum* and *Xeromphalina cornui* the abundance is relatively low but spatial frequency is high.

Many saprobic species present in the earlier succession phases attain the highest frequency and abundance in the forest plots, e.g., *Collybia cirrhata*, *C. tuberosa*, *Coniophora arida*, *Dacrymyces stillatus*, *Galerina pumila*, *Leptosporomyces galzinii*, *Mycena galopus*, *M. metata*, *M. zephirus*, *Phlebiella sulphurea* and *Xeromphalina cornui*. The number and frequency of carpophores in other species clearly decreases, for instance in *Clitocybe candidans*, *C. diatreta*, *Lycoperdon nigrescens*, *Macrolepiota procera*, *Mycena flavoalba* and *M. pura*. The most conspicuous "successional relict" is *Psilocybe montana*, a species very abundant in the optimal stage of vegetation development, typical of open psammophilous grassland with mosses and lichens. In the forest, only one carpophore was found.

Numerous saprobic species were found exclusively in phase 9/10 of the succession studied, e.g., *Athelia decipiens*, *Botryobasidium vagum*, *Crepidotus applanatus*, *Dacrymyces*



*ovisporus*, *D. macnabbii*, *Gymnopilus penetrans*, *Hypholoma capnoides*, *Mycena purpureofusca* and *Pseudomerulius aureus*. A great majority of them are wood-inhabiting fungi.

In nearly all subplots of the forest plots *Exobasidium vaccinii* was present, a leaf parasite of *Vaccinium vitis-idaea*.

## 5.2. Changes in species and sporocarp distribution along successional gradient

### 5.2.1. Species richness and diversity

During three years of observations 300 macrobasidiomycete species were found in permanent plots, among them 98 ECM (32.7%), 200 saprotrophic (66.7%) and 2 parasitic species (0.7%).

The lowest number of species (15) was found in phase 2 of the vegetation development (Fig. 9); all of them were saprobic fungi, and the majority of them were growing on herbaceous plant remnants. The number of species in the next phases gradually increases (with a slight decrease in phase 6/7) till the phase of juniper-aspen brushwood (phase 8), where 183 species were found. Although two thirds of them were saprotrophs and the proportion of the ECM species here was the lowest in the whole chronosequence, the actual number of these species was the highest. The highest proportion of the ECM fungi is characteristic of the community in the initial stage of succession (phase 4). Young, 6-8(12) year old trees, support a rich and diverse range of symbiotic species, while the group of saprotrophs is relatively poor. The highest numbers of the ECM species at plot level were recorded in phase 6 (38 species in each of the two plots).

The majority of the ECM species of the series studied are more or less host-specific to pine, e.g., *Suillus luteus*, *Chalciporus piperatus*, *Hygrophorus hypothejus*, *Inocybe lacera*, *Cortinarius cinnamomeoluteus*, *Lactarius rufus*, *Thelephora terrestris*, *Tricholoma albobrunneum*, *T. equestre* i *T. portentosum*. Most frequently under pine were also recorded species which could be associated with a broad range of both coniferous and deciduous trees, like *Amanita muscaria*, *Boletus edulis* and *Tricholoma saponaceum* var. *saponaceum*. Other non-specific fungi were found as frequently under pine as under birch or aspen, for instance *Paxillus involutus*, *Amanita rubescens*, *Laccaria proxima*, *Clitopilus prunulus* and *Amanita citrina*. The most common deciduous host-specific species were, e.g., *Leccinum aurantiacum* (aspen), *Russula fragilis* (birch, aspen), *R. clavipes* (birch, aspen) and *R. galochroa* (birch). Not surprisingly, this group is most numerous and diverse in juniper-aspen brushwood.

Saprotrophic fungi can be assigned to four main ecological groups: fungi growing on soil, litter-inhabiting fungi, fungi on dead wood and bryophilous species growing among mosses. Many of them were recorded not only on one but on different substrates.

Among the fungi most frequently found on soil were species of *Entoloma*, *Rhodocybe*, *Hygrocybe*, *Macrolepiota*, *Lycoperdon*, *Geastrum*, as well as *Hygroaster asterosporus* and *Ripartites tricholoma*. Most of them are typical of open areas like grasslands, heathlands or juniper shrubs.

The group of litter-inhabiting species were most diverse; they grew on decaying shoots of herbaceous plants (*Mycena olivaceomarginata*, *Parasola plicatilis*, *Crinipellis scabella*), fallen leaves of deciduous trees (*Gymnopus dryophilus*, *Lepista nuda*, *Clitocybe phyllophila*, *Marasmius epiphyllus*), juniper needles (*Clitocybe diatrete*, *C. fragrans*, *C. vibecina*), pine

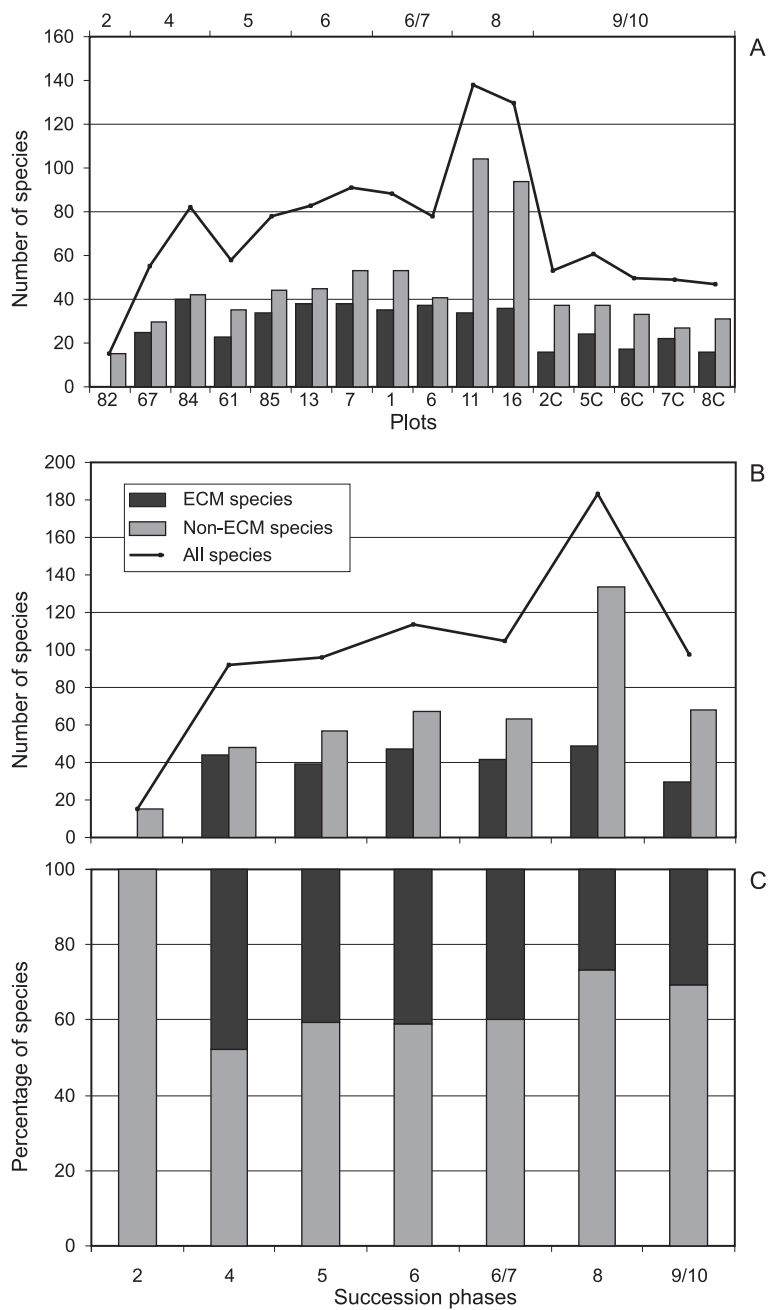


Fig. 9. Number of ECM and non-ECM species in plots (A) and succession phases (B) and percentage of these groups in the different succession phases (C).

needles (numerous species of *Mycena* and *Clitocybe*, *Gymnopus androsaceus*, *Xeromphalina cornui*), cones (*Baeospora myosura*, *Auriscalpium vulgare*, *Strobilurus stephanocystis*), twigs and pieces of bark (*Tubaria furfuracea*, *Flammulaster granulosus*, *Crucibulum laeve*, *Sphaerobolus stellatus*), as well as decaying fruit bodies of other fungi (*Collybia cirrhata*, *C. tuberosa*, *C. cookei*) and dung (*Panaeolus papilionaceus*).

Similarly, various fungi showed preferences for certain kinds of woody substrates: *Amylostereum laevigatum* grew on the wood of juniper, *Calocera viscosa*, *Gymnopilus penetrans*, *Phanerochaete sanguinea*, *Phlebiella sulphurea* – on the wood of pine, *Inonotus rheades*, *Daedaleopsis confragosa*, *Peniophora polygonia* – on the wood of aspen, *Stereum hirsutum*, *Fomes fomentarius* – on the wood of birch. Some species usually occurred on relatively freshly fallen branches, for example some species of *Hyphoderma*, *Peniophora* and *Stereum*, on branches and larger pieces of decorticated wood were species such as *Dacrymyces stillatus*, *Coniophora arida* and *Botryobasidium vagum* while on rotting logs and stumps were species like *Hypholoma* spp., *Pseudohydnum gelatinosum*, *Steccherinum fimbriatum* and *Mycena stipitata*. On dead branches of trees or on dead standing trees were found species like *Exidia plana*, *Tremella mesenterica*, *Dacrymyces variisporus*, *Inonotus rheades*, *Fomes fomentarius* and *Daedaleopsis confragosa*.

Many saprotrophic species did not show any special preferences and they occurred on wood in different stage of decay, on coarse as well as on fine debris, on wood and on litter and on different types of litter.

The bryophilous fungi are also a distinct ecological group. They include species of *Galerina* (*G. atkinsoniana*, *G. pumila*, *G. hypnorum*, *G. mniophila*, *G. calyptrata*, etc.), *Cystoderma* (*C. amiantinum*, *C. jasonis* var. *jasonis*, *C. jasonis* var. *lilacipes*) and *Cystodermella granulosa*, *Psilocybe montana*, *Rickenella fibula*, *Cantharellula umbonata*. The last species is a moss parasite and occurred in *Polytrichum* tufts, mainly on *P. juniperinum* and *P. commune*.

Another parasitic fungus collected was *Exobasidium vaccinii* growing abundantly on leaves of *Vaccinium vitis-idaea*.

The macrofungal species diversity changes along with the succession process (Fig. 10). For the non-ECM fungi, changes in diversity index reflect the increasing number of species in subsequent phases of succession (Fig. 9; *Dacrymyces stillatus*, the species producing huge

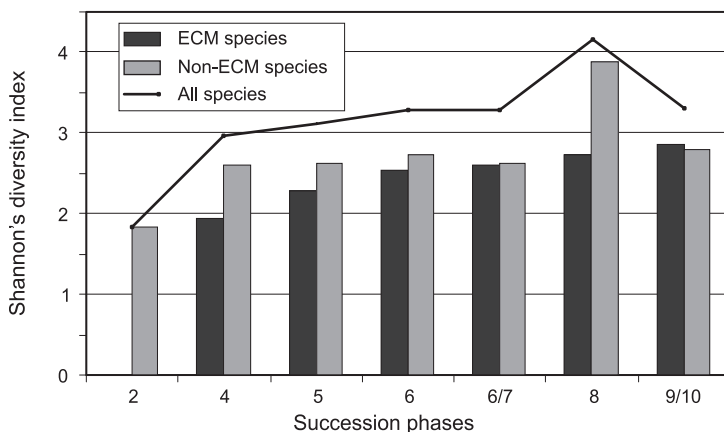


Fig. 10. Species diversity of fungi in consecutive phases of the spontaneous secondary succession on abandoned farmland.



Phases	2	4	5	6	6/7	8	9/10
2		25.4	27.8	24.4	25.6	10.7	12.1
4	—		72.4	60.9	57.7	35.2	31.0
5	—	69.9		61.3	63.3	35.6	32.0
6	—	63.7	67.4		67.7	41.8	35.6
6/7	—	62.8	66.7	87.6		45.7	47.3
8	—	62.4	56.8	54.2	50.6		41.6
9/10	—	32.4	26.1	28.6	27.8	32.9	

ECM species

Non-ECM species

Fig. 11. Diagram showing Sørensen's similarity index of fungal species among consecutive phases of the spontaneous secondary succession on abandoned farmland.

numbers of very tiny carpophores, was excluded from calculations); the diversity is the highest in phase 8, reaching 3.88 at phase level (calculation based on the data from two plots taken together, i.e. 2000 m<sup>2</sup>) and 3.54 and 3.66 at plot level (calculation based on the data from the two separate plots of 1000 m<sup>2</sup>; data not shown). For the ECM fungi, however, diversity increases relatively independently of species number fluctuations, which is an effect of higher species evenness (the Shannon's diversity index is increased either by having additional species, or by having a greater species evenness). It is particularly clear in the terminal phase of succession, in pine forest, where the ECM fungi attain the highest diversity at phase level – 2.87 despite the relatively low number of species (Fig. 10). At plot level, ECM fungal diversity grows rapidly up to 2.55 and 2.59 in phase 6/7, then slightly decreases and increases again in the pine forest (data not shown).

The species composition of fungi is very similar in the plots representing the same phase of succession (based on Sørensen's index; see chapter 5.1.), from 60.9% to 83.3%. High similarity, usually exceeding 60%, is also a rule in the assemblages of fungi occurring in phases 4-6/7 (late initial and optimal stage of vegetation succession) (Fig. 11). The production of carpophores of the ECM species, which starts between phases 2 and 4 in the initial stage of succession, undergoes mainly quantitative changes in subsequent pre-forest phases. A distinct qualitative change can be seen between phases 8 and 9/10 when after the transitional community of juniper-aspen brushwood with pine the pine forest of *Peucedano-Pinetum* type is formed. Many species which occurred in earlier communities finish producing their carpophores at this point. In the pine forest, only about 50% of the ECM species, common with the earlier phases, still fruit and the similarity index decreases to about 30%.

A fundamental change in species composition of saprotrophic fungi takes place twice. The first time occurs between phases 2 and 4, when dense herb-grass vegetation with fungi inhabiting abundant necromass of herbaceous plants is replaced by poor, psammophilous grassland where fungi growing on soil predominate as well as the bryophilous species and fungi growing on litter accumulating under pines and junipers. The second time is between phases 6/7 and 8, when the canopies of trees close and accumulation of litter and woody debris increases. Relatively low species similarity of saprobic fungi between juniper-aspen brushwood and the earlier vegetation types results mainly from the great number of the brushwood-exclusive species inhabiting litter and woody debris originating from deciduous trees. The species similarity between phases 4-6 and 9/10 is low because of the arrival of numerous species growing on decaying pine wood.

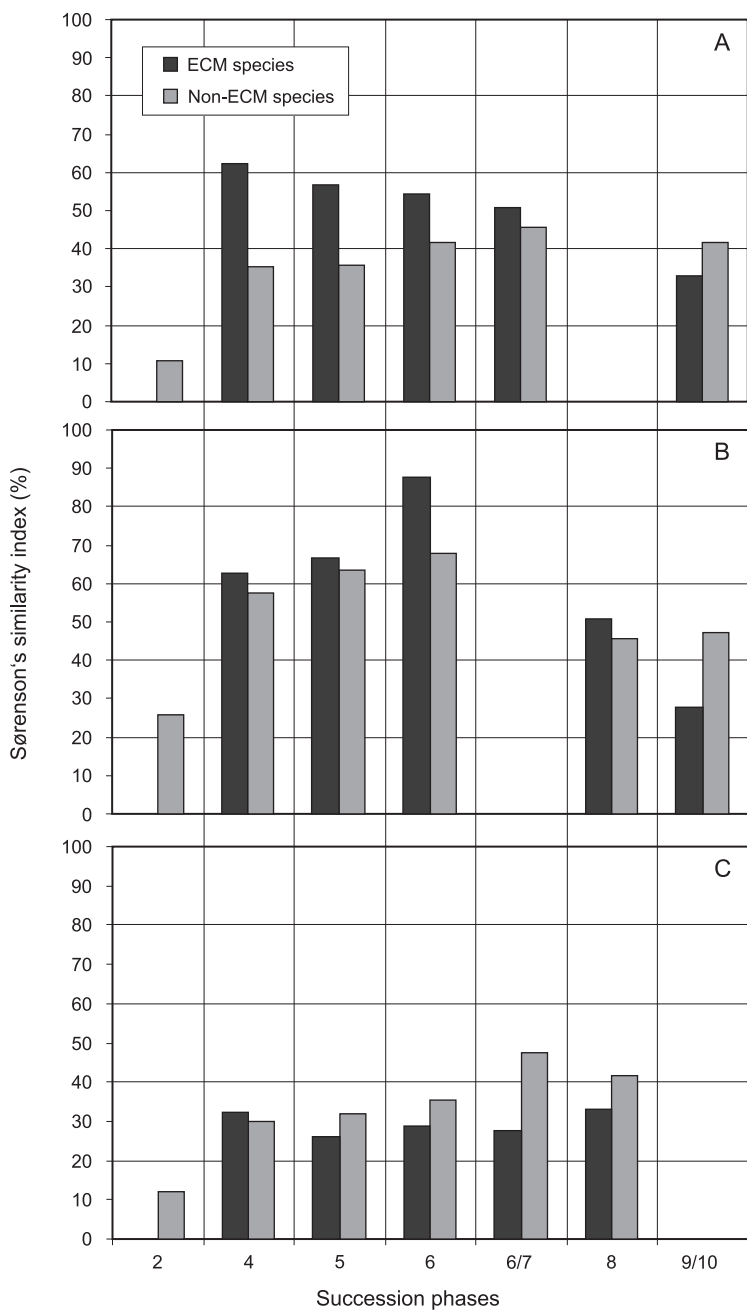


Fig. 12. Similarity in species composition among phases 6/7 (A), 8 (B) and 9/10 (C) and the other succession phases.

Sørensen's index of fungal species similarity between the succession phases is usually higher for the communities which are closer in the chronosequence (Fig. 12, see also Fig. 11) with an exception of phase 8, where the species composition of the ECM fungi is most similar to phase 4. This results from the higher proportion of deciduous trees in the plots representing phases 4 and 5 than in the plots of phases 6 and 6/7 (see Appendix 2, Tab. A2.1). Moreover, both plots of phase 4 and one plot of phase 5 are situated in the immediate neighbourhood of juniper-aspen brushwood (phase 8).

Most of the typical forest ECM species arrive only in the final phase of the chronosequence studied; this is reflected by the relatively low and stable similarity between the ECM species composition of pine forest (phase 9/10) and pre-forest vegetation.

### 5.2.2. Species abundance patterns

Fungal populations change with time and vegetation development, showing specific patterns of sporocarp production which can be seen in abundance changes in subsequent succession phases as well as in changes of spatial distribution. Groupings of fungal species showing similar dynamics along successional gradient were revealed with the k-mean analysis based on density of carpophores [ $y = \ln(\text{DC}3y + 1)$ ] (Appendix 4).

#### ECM fungi

For the ECM fungi five species clusters were distinguished (Fig. 13; Appendix 4, Tab. A4.1):

**Cluster 1.** Species producing large numbers of carpophores in the initial and optimal stage of succession, and considerably fewer carpophores in the terminal stage. These are: *Amanita muscaria*, *Chalciporus piperatus*, *Cortinarius cinnamomeoluteus*, *Hygrophorus hypothecus*, *Inocybe lacera*, *Laccaria proxima* and *Suillus luteus*.

**Cluster 2.** Species producing high numbers of carpophores in the optimal stage of vegetation succession, especially in its advanced phases, and low (or lower) numbers in the other stages. Examples include: *Boletus edulis*, *Cortinarius parevernius*, *Laccaria bicolor*, *Lactarius rufus*, *Rhodocollybia butyracea* f. *asema*, *Thelephora terrestris*, *Tricholoma albobrunneum*, *T. equestre*, *T. portentosum* and *T. saponaceum* var. *saponaceum*.

**Cluster 3.** Species specific to or attaining the highest abundance in transitional vegetation type of juniper-aspen brushwood, e.g., *Cantharellus cibarius*, *Clitopilus prunulus*, *Cortinarius albobviolaceus*, *C. trivialis*, *Inocybe flocculosa*, *Lactarius torminosus*, *Leccinum aurantiacum*, *Russula clavipes*, *R. galochroa* and *Thelephora caryophyllea*.

**Cluster 4.** Species showing preference towards forest community, most of them arriving only in well developed pine forest; in the open vegetation they usually occur sporadically and produce few carpophores while in the forest their abundance is considerably higher. These are *Amanita porphyria*, *Cortinarius caperatus*, *C. semisanquineus*, *C. vibratilis*, *Paxillus involutus*, *Russula badia*, *R. decolorans*, *R. emetica*, *R. paludosa* and *Xerocomus badius*.

**Cluster 5.** Species producing relatively low numbers of carpophores, the majority of them showing less strong differences in carpophore production in particular succession phases as compared with the species grouped in the former clusters. Over half of the ECM species

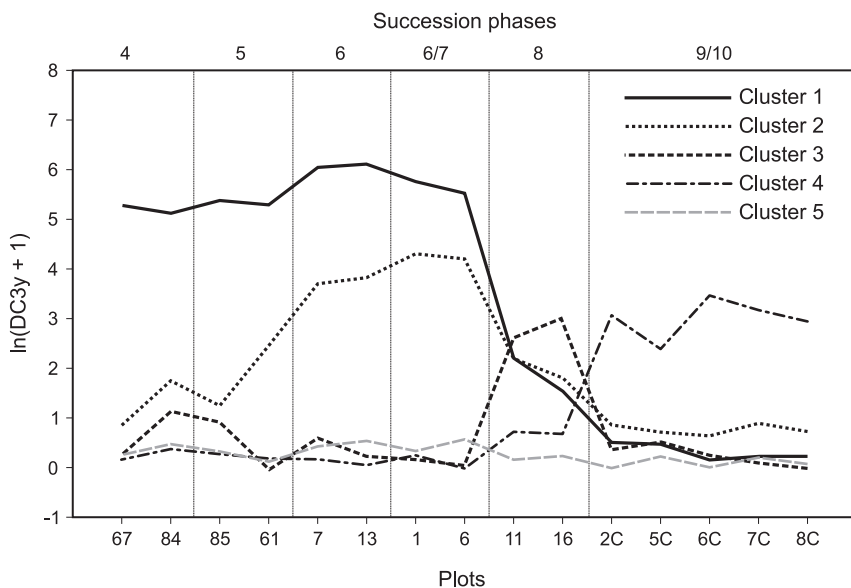


Fig. 13. ECM species grouping according to k-mean analysis ( $k = 4$ ). For species list in each cluster and distance from k-mean see Appendix 4.

recorded belong to this cluster (54). They include fungi occurring in a few different succession phases or just in one succession stage, as well as numerous exclusive species found in the plots representing the same succession phase or just in a single plot.

Among the ECM fungi there is a distinct but not numerous group of species producing sporocarps in all the succession phases, starting from (at least) phase 4 (Tab. 1). *Amanita rubescens* seems to occur relatively independently of vegetation development, while the other species show preferences towards some of the succession phases. *Laccaria proxima* and most of the species from the *Cortinarius* spp. I group (with small carpophores) occur mainly in open vegetation of initial and optimal succession stage and they tend to disappear in forest community (cluster 1). *Thelephora terrestris* and most of the species from the groups *Cortinarius* spp. II and spp. III (medium-size carpophores) as well as *Rhodocollybia butyracea* f. *asema* occur mainly in advanced phases of the optimal stage (cluster 2), the latter species continuing abundant fruiting also in transitional pre-forest community. *Russula fragilis* appears independently of vegetation development but always associated with deciduous trees and most abundantly in juniper-aspen brushwood (cluster 3). *Paxillus involutus*, rather scattered in all open communities, clearly prefers the forest phase (cluster 4).

A large group of the ECM species are associated with pre-forest vegetation type, i.e. from psammophilous grassland with single trees to transitional juniper-aspen brushwood (Tab. 1). This is especially true for the fungi grouped in clusters 1 and 2. Most of them seem to have their ecological optimum in particular succession phases or stages: *Inocybe lacera* in the initial stage (Fig. 14), *Suillus luteus* in the initial and optimal stage, *Cortinarius cinnamomeoluteus* in the early phase of the optimal stage, *Hygrophorus hypothejus* and *Amanita muscaria* in the optimal stage (Fig. 15), *Tricholoma albobrunneum*, *T. equestre* and *T. portentosum* in the last phase of optimal stage (Fig. 16). These species produce carpophores also in the juniper-aspen brushwood but, as a rule, less abundantly.

Table 1  
Distribution of the ECM species in the old-field chronosequence

Parameter	aDCy							aSF						
Succession phase	2	4	5	6	6/7	8	9/10	2	4	5	6	6/7	8	9/10
<i>Laccaria proxima</i>	·	102.7	149.2	192.2	171.3	1.7	1.3	·	VI	VI	VII	VII	II	III
<i>Cortinarius</i> spp. I	·	81.8	74.8	439.0	282.7	41.3	4.5	·	VI	V	IX	X	VIII	III
<i>Thelephora terrestris</i>	·	8.0	13.0	58.7	21.8	0.2	12.0	·	I	III	V	III	I	IV
<i>Cortinarius</i> spp. II	·	14.3	7.2	3.5	11.8	97.7	3.2	·	II	II	I	II	X	III
<i>Rhodocollybia butyracea</i> f. <i>asema</i>	·	0.7	4.3	17.3	21.7	33.3	1.0	·	I	I	IV	IV	IX	I
<i>Russula fragilis</i>	·	11.0	6.5	9.0		12.5	5.8	·	II	II	I		VII	II
<i>Cortinarius</i> spp. III	·	8.8	25.3	3.7	1.7	22.8	1.2	·	I	II	II	I	V	II
<i>Paxillus involutus</i>	·	1.5	4.7	0.3	0.3	6.2	25.0	·	I	I	I	I	II	IX
<i>Amanita rubescens</i>	·	1.8	3.8	0.8	1.3	0.3	0.7	·	I	I	I	I	I	II
<i>Tricholoma saponaceum</i> var. <i>squamosum</i>	·	4.2	·	·	·	·	2.0	·	I	·	·	·	·	I
<i>Inocybe lacera</i>	·	538.3	195.8	43.2	24.0	0.3	·	·	IX	V	V	III	I	·
<i>Suillus luteus</i>	·	49.7	82.0	93.0	54.5	2.8	·	·	VIII	IX	IX	VII	II	·
<i>Cortinarius cinnamomeoluteus</i>	·	137.8	281.7	288.3	321.5	0.5	·	·	V	VIII	X	X	I	·
<i>Hygrophorus hypothejus</i>	·	23.8	116.0	363.5	186.3	0.8	·	·	V	VIII	X	IX	I	·
<i>Amanita muscaria</i>	·	76.3	70.5	154.3	121.3	18.2	·	·	VII	VII	IX	X	VIII	·
<i>Chalciporus piperatus</i>	·	19.7	9.2	63.7	27.3	1.3	·	·	IV	IV	VIII	VII	III	·
<i>Tricholoma albobrunneum</i>	·	0.3	19.8	50.3	73.3	0.7	·	·	I	III	VII	IX	I	·
<i>Tricholoma portentosum</i>	·	2.3	3.5	43.5	84.5	3.0	·	·	I	I	V	VII	III	·
<i>Tricholoma equestre</i>	·	·	3.2	27.5	82.8	13.8	·	·	·	I	IV	VIII	VIII	·
<i>Leccinum aurantiacum</i>	·	3.2	3.7	3.5	0.2	8.0	·	·	II	II	I	I	IV	·
<i>Boletus edulis</i>	·	1.7	3.0	1.3	6.8	1.7	·	·	I	I	I	III	III	·
<i>Clitopilus prunulus</i>	·	1.8	3.7	5.5	1.7	4.5	·	·	I	I	I	I	III	·
<i>Amanita pantherina</i>	·	1.5	0.3	0.7	0.7	0.7	·	·	I	I	I	I	I	·
<i>Hebeloma</i> sp. 1	·	1.2	·	2.0	·	0.5	·	·	I	·	I	·	I	·
<i>Russula galochroa</i>	·	1.8	0.2	·	·	3.8	·	·	I	I	·	·	V	·
<i>Thelephora caryophyllea</i>	·	1.0	·	·	·	2.7	·	·	I	·	·	·	II	·
<i>Hebeloma</i> sp. 2	·	1.5	·	·	·	0.3	·	·	I	·	·	·	I	·
<i>Coltricia perennis</i>	·	0.5	·	·	·	0.2	·	·	I	·	·	·	I	·
<i>Leccinum scabrum</i>	·	0.3	·	·	·	1.0	·	·	I	·	·	·	II	·
<i>Inocybe calida</i>	·	1.7	0.5	0.3	0.2	·	·	·	I	I	I	I	·	·
<i>Inocybe praetervisa</i>	·	0.5	0.7	0.5	0.2	·	·	·	I	I	I	I	·	·
<i>Laccaria bicolor</i>	·	3.8	0.3	37.8	16.2	·	·	·	I	I	II	II	·	·
<i>Ramaria eumorpha</i>	·	0.5	·	14.5	0.8	·	·	·	I	·	I	I	·	·
<i>Cortinarius purpurascens</i>	·	2.3	·	1.2	1.8	·	·	·	I	·	I	II	·	·
<i>Inocybe assimilata</i>	·	0.3	·	1.2	4.7	·	·	·	I	·	I	I	·	·
<i>Boletus reticulatus</i>	·	0.3	·	·	0.3	·	·	·	I	·	·	I	·	·
<i>Inocybe curvipes</i>	·	·	2.0	3.3	0.7	·	·	·	·	I	I	I	·	·
<i>Cortinarius anthracinus</i>	·	·	0.5	5.0	7.5	·	·	·	·	I	II	III	·	·
<i>Tricholoma saponaceum</i> var. <i>saponaceum</i>	·	·	5.3	23.5	44.5	·	·	·	·	I	II	IV	·	·
<i>Lactarius rufus</i>	·	·	·	29.2	104.7	0.5	12.7	·	·	·	IV	V	I	VI
<i>Cortinarius parevernus</i>	·	·	·	26.8	20.0	0.3	·	·	·	·	IV	IV	I	·
<i>Inocybe asterospora</i>	·	9.2	·	0.2	·	·	·	·	II	·	I	·	·	·
<i>Hygrophorus aureus</i>	·	0.7	·	·	·	·	·	·	I	·	·	·	·	·
<i>Lactarius vietus</i>	·	0.3	·	·	·	·	·	·	I	·	·	·	·	·
<i>Xerocomus subtomentosus</i>	·	1.0	0.8	·	·	·	·	·	I	I	·	·	·	·
<i>Russula subfetens</i>	·	0.3	0.2	·	·	·	·	·	I	I	·	·	·	·
<i>Gyroporus castaneus</i>	·	0.5	0.5	·	·	·	·	·	I	I	·	·	·	·
<i>Lactarius quieticolor</i>	·	·	2.8	·	·	·	·	·	·	I	·	·	·	·



<i>Scleroderma fuscum</i>	.	.	0.5	.	.	.	.	.	.	I	.	.	.	.
<i>Xerocomus chrysenteron</i>	.	.	0.3	.	.	.	.	.	.	I	.	.	.	.
<i>Russula vinosobrunnea</i>	.	.	0.2	.	.	.	.	.	.	I	.	.	.	.
<i>Cortinarius cinnabarinus</i>	.	.	0.7	0.2	.	.	.	.	.	I	I	.	.	.
<i>Cortinarius colus</i>	.	.	.	9.3	.	.	.	.	.	.	II	.	.	.
<i>Tricholoma pessundatum</i>	.	.	.	0.5	.	.	.	.	.	.	I	.	.	.
<i>Cortinarius acetosus</i>	.	.	.	0.2	.	.	.	.	.	.	I	.	.	.
<i>Cortinarius mucosus</i>	.	.	.	9.0	4.0	.	.	.	.	.	III	II	.	.
<i>Cortinarius multififormis</i> var. <i>coniferarum</i>	.	.	.	3.5	0.7	.	.	.	.	.	I	I	.	.
<i>Rhizopogon luteolus</i>	.	.	.	0.3	0.3	.	.	.	.	.	I	I	.	.
<i>Suillus bovinus</i>	.	.	.	0.3	0.8	.	.	.	.	.	I	I	.	.
<i>Sarcodon squamosus</i>	.	.	.	2.2	5.0	.	.	.	.	.	I	II	.	.
<i>Cortinarius</i> spp. IV	.	.	.	0.2	2.8	.	.	.	.	.	I	I	.	.
<i>Cortinarius sanguineus</i>	.	.	.	.	1.0	.	.	.	.	.	.	I	.	.
<i>Boletus pinophilus</i>	.	.	.	.	0.3	.	.	.	.	.	.	I	.	.
<i>Russula clavipes</i>	.	0.3	0.2	.	.	10.3	.	.	I	I	.	.	VIII	.
<i>Lactarius uvidus</i>	.	.	0.2	.	.	3.0	.	.	.	I	.	.	II	.
<i>Tricholoma imbricatum</i>	.	.	.	0.7	.	3.2	.	.	.	.	I	.	I	.
<i>Cortinarius trivialis</i>	.	.	.	.	.	117.3	.	.	.	.	.	.	X	.
<i>Cortinarius alboviolaceus</i>	.	.	.	.	.	17.5	.	.	.	.	.	.	V	.
<i>Hebeloma</i> sp. 3	.	.	.	.	.	16.3	.	.	.	.	.	.	IV	.
<i>Lactarius torminosus</i>	.	.	.	.	.	4.3	.	.	.	.	.	.	IV	.
<i>Inocybe flocculosa</i>	.	.	.	.	.	11.5	.	.	.	.	.	.	III	.
<i>Inocybe</i> cf. <i>margaritispota</i>	.	.	.	.	.	3.0	.	.	.	.	.	.	III	.
<i>Amanita gemmata</i>	.	.	.	.	.	0.8	.	.	.	.	.	.	II	.
<i>Lactarius necator</i>	.	.	.	.	.	0.5	.	.	.	.	.	.	I	.
<i>Ramaria</i> cf. <i>abietina</i>	.	.	.	.	.	0.5	.	.	.	.	.	.	I	.
<i>Russula mustelina</i>	.	.	.	.	.	0.5	.	.	.	.	.	.	I	.
<i>Amanita phalloides</i>	.	.	.	.	.	0.2	.	.	.	.	.	.	I	.
<i>Hebeloma stenocystis</i>	.	.	.	.	.	0.2	.	.	.	.	.	.	I	.
<i>Russula versicolor</i>	.	.	.	.	.	0.2	.	.	.	.	.	.	I	.
<i>Amanita citrina</i>	.	0.5	.	.	.	39.0	19.8	.	I	.	.	.	X	IX
<i>Cantharellus cibarius</i>	.	.	.	.	.	22.8	2.5	.	.	.	.	.	III	III
<i>Russula paludosa</i>	.	.	.	.	.	0.8	6.5	.	.	.	.	.	I	VII
<i>Russula emetica</i>	.	0.5	.	.	.	.	13.2	.	I	.	.	.	.	VII
<i>Xerocomus badius</i>	.	.	.	0.5	0.8	.	8.7	.	.	I	I	.	.	VI
<i>Cortinarius caperatus</i>	.	.	.	.	.	.	15.3	.	.	.	.	.	.	VIII
<i>Russula decolorans</i>	.	.	.	.	.	.	10.2	.	.	.	.	.	.	VIII
<i>Cortinarius semisanguineus</i>	.	.	.	.	.	.	16.3	.	.	.	.	.	.	V
<i>Russula badia</i>	.	.	.	.	.	.	9.2	.	.	.	.	.	.	V
<i>Amanita porphyria</i>	.	.	.	.	.	.	2.8	.	.	.	.	.	.	IV
<i>Cortinarius vibratilis</i>	.	.	.	.	.	.	2.8	.	.	.	.	.	.	III
<i>Russula xerampelina</i>	.	.	.	.	.	.	1.3	.	.	.	.	.	.	III
<i>Russula vinosa</i>	.	.	.	.	.	.	1.7	.	.	.	.	.	.	II
<i>Lactarius tabidus</i>	.	.	.	.	.	.	2.0	.	.	.	.	.	.	I
<i>Russula turci</i>	.	.	.	.	.	.	0.8	.	.	.	.	.	.	I
<i>Chroogomphus rutilus</i>	.	.	.	.	.	.	0.2	.	.	.	.	.	.	I
<i>Russula vesca</i>	.	.	.	.	.	.	0.2	.	.	.	.	.	.	I
<i>Suillus variegatus</i>	.	.	.	.	.	.	0.2	.	.	.	.	.	.	I
<i>Tylopilus felleus</i>	.	.	.	.	.	.	0.2	.	.	.	.	.	.	I

aDCy – average density of carpophores per year; aSF – average spatial frequency expressed as classes: I=1-10%, II=11-20%, III=21-30%, IV=31-40%, V=41-50%, VI=51-60%, VII=61-70%, VIII=71-80%, IX=81-90%, X=91-100% (the transformation makes the data comparable as the numbers of subplots in succession phases are different: phase 2 – 10, phase 4 – 80, phase 5 – 80, phase 6 – 80, phase 6/7 – 80, phase 8 – 33, phase 9/10 – 20)

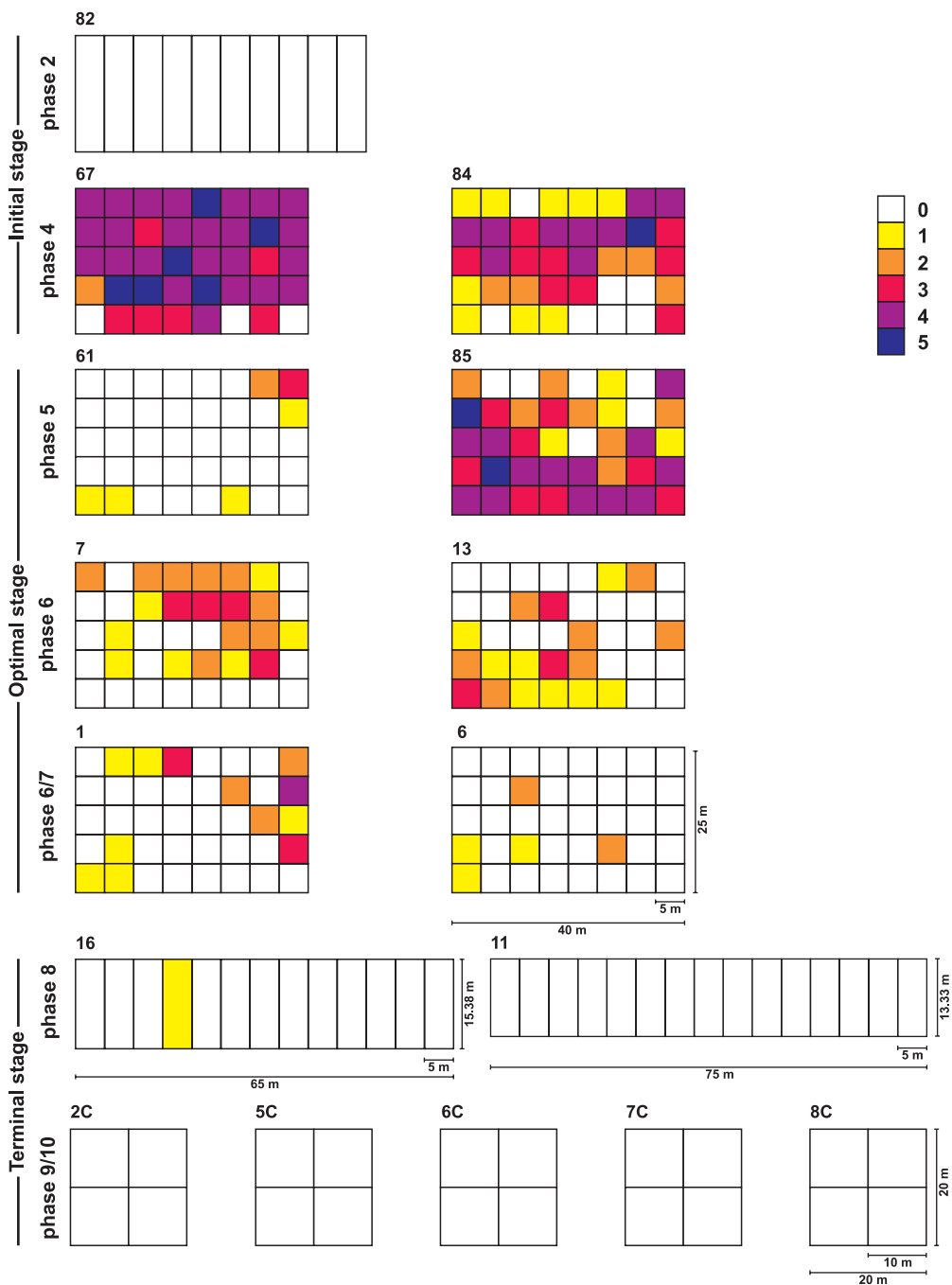


Fig. 14. *Inocybe lacera* – distribution in the plots based on the total abundance of carpophores (AC3y) in each subplot according to the scale by KEIZER (1993a): 1 = 1-3, 2 = 4-11, 3 = 12-32, 4 = 33-89, 5 = 90-243 carpophores.

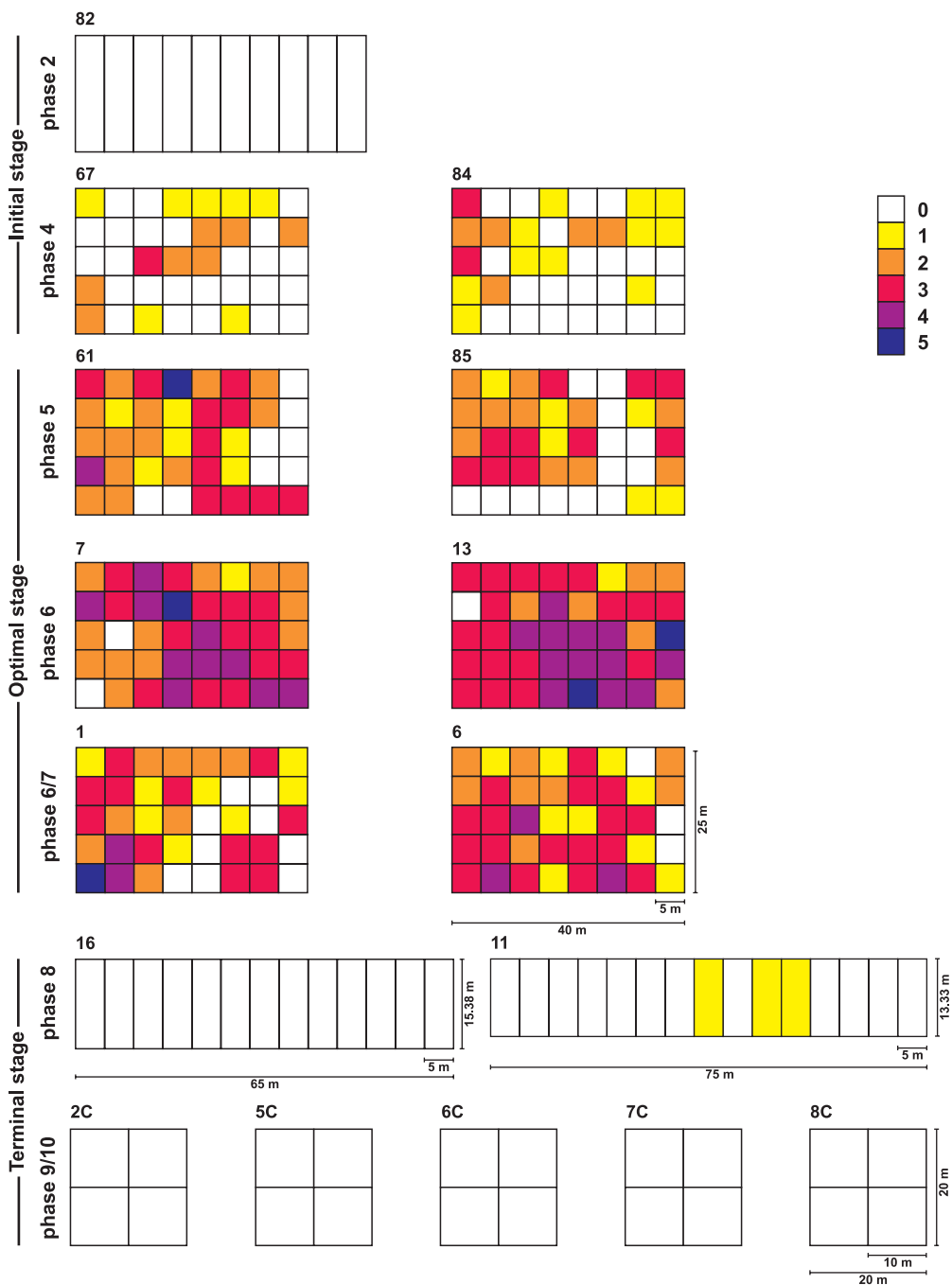


Fig. 15. *Hygrophorus hypothejus* – distribution in the plots based on the total abundance of carpophores (AC3y) in each subplot according to the scale by KEIZER (1993a): 1 = 1-3, 2 = 4-11, 3 = 12-32, 4 = 33-89, 5 = 90-243 carpophores.

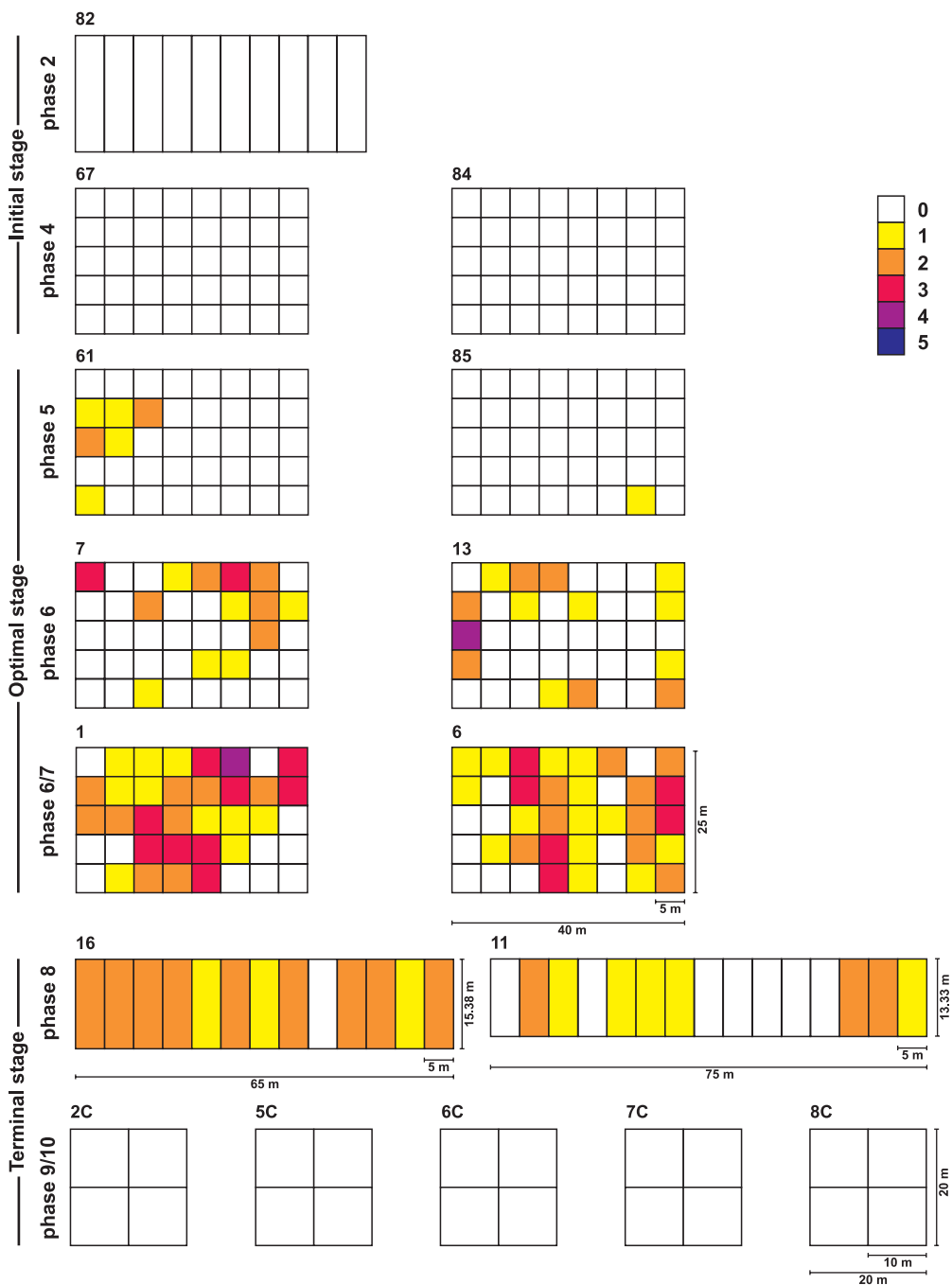


Fig. 16. *Tricholoma equestre* – distribution in the plots based on the total abundance of carpophores (AC3y) in each subplot according to the scale by KEIZER (1993a): 1 = 1-3, 2 = 4-11, 3 = 12-32, 4 = 33-89, 5 = 90-243 carpophores.

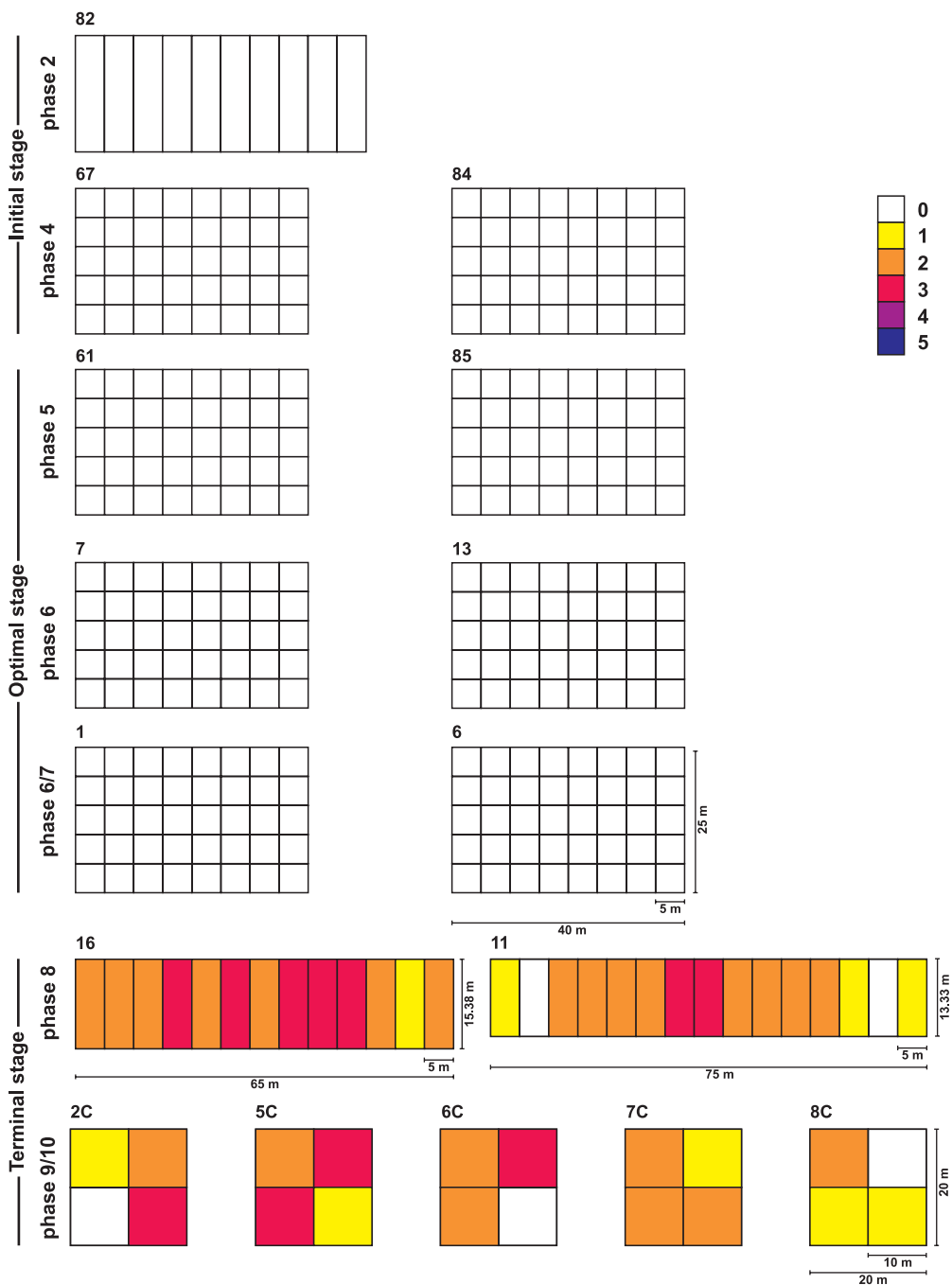


Fig. 17. *Amanita citrina* – distribution in the plots based on the total abundance of carpophores (AC3y) in each subplot according to the scale by KEIZER (1993a): 1 = 1-3, 2 = 4-11, 3 = 12-32, 4 = 33-89, 5 = 90-243 carpophores.



Among the species which clearly prefer open habitats and do not appear (as sporocarps) in brushwood community are most of the *Inocybe* species, *Laccaria bicolor*, *Cortinarius purpurascens* and *C. anthracinus*.

The majority of the ECM species attaining the highest abundance in the juniper-aspen brushwood (phase 8, cluster 3) are the symbionts of deciduous trees. Some of them, like *Leccinum aurantiacum*, *L. scabrum*, *Russula galochroa* and *R. clavipes*, accompany aspen and birch also in the plots of earlier succession phases, especially in places directly neighbouring the brushwood. Other species such as *Cortinarius alboviolaceus*, *C. trivialis*, *Inocybe flocculosa* and *Lactarius torminosus* occur exclusively in phase 8.

Most of the ECM fungi that occur in a few different succession phases arrive and start fruiting relatively early, in the initial stage or in the early phase of optimal stage, and they cease production of their carpophores in late pre-forest communities of phases 6/7 or 8 (Tab.1). The species that occur both in pre-forest and in forest sites are relatively few; apart from the several species listed earlier in this chapter that occur in the chronosequence at least from phase 4; among such species are *Russula emetica* and *Xerocomus badius* – occurring sporadically out of forest (cluster 4), *Amanita citrina*, *Cantharellus cibarius* and *Russula paludosa* – not occurring in open vegetation (clusters 3 and 4) (Fig. 17) and *Lactarius rufus* – associated mainly with younger coniferous stands (cluster 2). Almost half of the species found in the forest do not occur earlier in the chronosequence. Some of them produce relatively abundant carpophores, for example *Amanita porphyria*, *Cortinarius caeperatus*, *C. semisanguineus*, *C. vibratilis*, *Russula badia* and *R. decolorans* (cluster 4).

The number of ECM exclusive species is low in the initial and optimal stages of the vegetation succession (Tab.1) and these are sporadic species attaining low abundance and spatial frequency. In the terminal stage, the number of exclusive species distinctly increases and many of them attain high abundance and high frequency.

### Non-ECM fungi

For the non-ECM fungi again five species clusters were distinguished (Fig. 18; Appendix 4, Tab. A4.2):

**Cluster 1.** Species producing high numbers of carpophores in all the succession stages, from the initial communities to the pine forest. These include species of *Entoloma* and *Galerina* as well as *Clitocybe diatreta*, *Cystoderma amianthinum*, *Gymnopus androsaceus* and *Collybia cirrhata*.

**Cluster 2.** Species producing very abundant carpophores in the complex of psammophilous grassland, juniper shrubs and pine (phases 4-6/7) and usually less abundant outside these communities. These are *Ampulloclitocybe clavipes*, *Cantharellula umbonata*, *Clavaria argillacea*, *Galerina atkinsoniana*, *Hygroaster asterosporus*, *Lycoperdon perlatum*, *Psilocybe montana*, *Rickenella fibula*; all these species grow mainly among mosses and on soil. In this group are also *Mycena flavoalba*, *M. leptcephala* and *M. olivaceomarginata* fruiting most abundantly in phase 2 on the remnants of herbaceous plants.

**Cluster 3.** A well distinguished group of 47 species with optimum of occurrence in the juniper-aspen brushwood (phase 8), characterized by abundant carpophore production in this type of vegetation. The cluster includes both species found also in the other succession phases, such as *Clitocybe anisata*, *C. candicans*, *C. fragrans*, *Dacrymyces variisporus*, *Gymnopus confluentis*, *Lycoperdon molle*, *Macrolepiota procera*, *Mycena stylobates*, *Sphaerobolus*

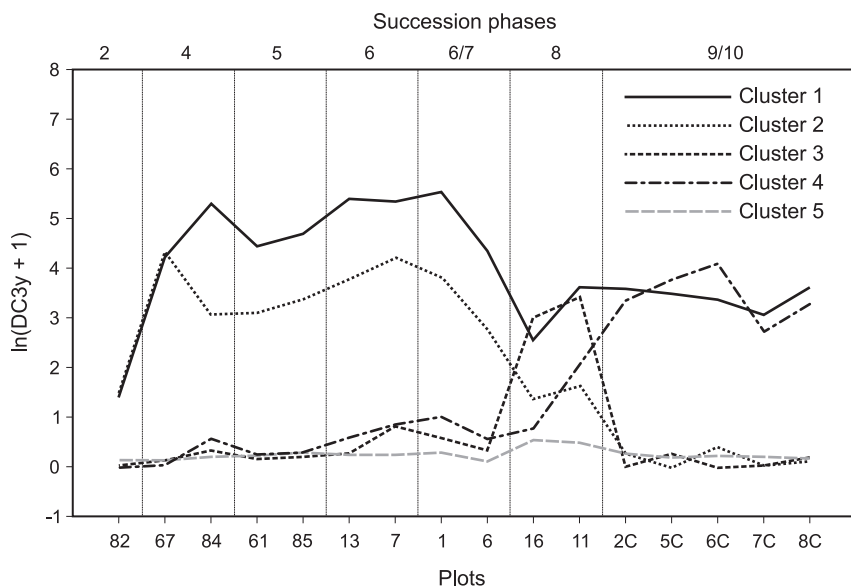


Fig. 18. Non-ECM species grouping according to k-mean analysis ( $k = 4$ ). For species list in each cluster and distance from k-mean see Appendix 4.

*stellatus*, and numerous exclusive species, such as *Clitocybe metachroa*, *Coprinellus* spp., *Crepidotus lundelii*, *Exidia plana*, *Flammulaster rhombosporus*, *Hymenochaete tabacina*, *Peniophora polygonia*, *Stereum hirsutum*.

**Cluster 4.** Species producing low numbers of carpophores in pre-forest vegetation, with optimum of occurrence and abundant carpophore production in the terminal stage of succession, particularly in the pine forest. Among them are typical litter-inhabiting species, e.g., *Auriscalpium vulgare*, *Baeospora myosura*, *Collybia tuberosa*, *Mycena galopus*, *M. metata*, *M. vulgaris*, *M. zephirus*, *Xeromphalina cornui*; wood-inhabiting species, e.g., *Dacrymyces stillatus*, *Hypholoma capnoides*, *Leptosporomyces galzinii*, *Phlebiella sulphurea*; as well as a parasite of *Vaccinium vitis-idaea*: *Exobasidium vaccinii*.

**Cluster 5.** Species producing relatively sparse carpophores, occurring in a few succession phases, and a majority of exclusive species. It is the most numerous group of species – over 120 taxa.

Most of the species which already appear in the initial stage of succession (phases 2 and 4) also occur in the optimal and terminal stage (Tab. 2). A great majority of them produce their carpophores much more abundantly in a particular succession phase or stage, depending on the type and amount of substrate available. For example, *Crinipellis scabella*, *Marasmius oreades*, *Mycena flavoalba*, *M. leptcephala*, *M. olivaceomarginata* prefer the habitats rich in the remnants of herbaceous plants (phase 2). *Clitocybe vibecina* and *Gymnopus androsaceus* occur abundantly on the sites of accumulation of row pine and juniper needles (late optimal and terminal stage). *Hypholoma fasciculare* and *Dacrymyces stillatus* are most numerous in the pine forest, where coniferous woody debris is most abundant.

Distinct differences in carpophore numbers produced in the subsequent succession phases can be seen mainly in the species growing on soil and among mosses, grouped in

Table 2  
Distribution of the non-ECM species in the old-field chronosequence

Parameter	aDCy								aSF							
	2	4	5	6	6/7	8	9/10		2	4	5	6	6/7	8	9/10	
<i>Crinipellis scabella</i>	227.5	·	5.3	3.0	·	·	·	X	·	I	I	·	·	·	·	
<i>Entoloma undatum</i>	0.5	·	·	·	0.2	·	·	I	·	·	·	I	·	·	·	
<i>Rickenella fibula</i>	3.0	20.2	24.3	110.3	12.2	·	·	I	V	IV	VI	II	·	·	·	
<i>Marasmius oreades</i>	113.0	·	0.2	1.7	1.2	2.0	·	X	·	I	I	I	I	I	·	
<i>Mycena olivaceomarginata</i>	56.0	12.7	3.2	1.3	1.5	7.5	·	IX	V	I	I	I	I	VI	·	
<i>Mycena leptocephala</i>	82.0	11.5	17.8	46.5	27.5	20.5	·	VIII	II	III	V	V	VIII	·	·	
<i>Mycena flavoalba</i>	104.5	30.7	14.8	4.7	5.5	21.5	0.2	X	II	II	I	II	V	I	·	
<i>Galerina</i> spp.	25.0	39.8	20.2	71.3	59.7	3.8	5.3	VII	IV	III	V	IV	III	IV	·	
<i>Clitocybe diatreta</i>	4.0	73.7	150.5	357.2	269.8	11.0	2.2	I	VIII	VII	IX	X	IV	I	·	
<i>Cystoderma amianthinum</i>	4.5	45.5	17.5	100.5	26.3	1.3	9.7	I	V	II	VII	IV	I	VII	·	
<i>Mycena pura</i>	3.0	1.2	0.8	3.7	1.3	17.8	0.5	IV	I	I	I	I	VIII	I	·	
<i>Psilocybe montana</i>	·	123.0	117.7	190.3	135.5	1.7	0.2	·	IX	IX	IX	IX	I	I	·	
<i>Entoloma</i> spp.	·	68.7	37.5	10.8	8.3	7.5	4.7	·	VIII	VI	IV	II	IV	V	·	
<i>Ampulloclitocybe clavipes</i>	·	2.7	0.5	20.3	5.3	8.0	2.2	·	I	I	III	II	III	III	·	
<i>Clitocybe vibecina</i>	·	5.8	1.3	10.7	45.7	1.0	12.3	·	I	I	II	III	II	IV	·	
<i>Gymnopus androsaceus</i>	·	54.2	21.0	120.0	204.5	77.7	159.0	·	II	I	V	V	V	X	·	
<i>Collybia cirrhata</i>	·	29.8	86.3	43.8	45.7	57.3	140.2	·	III	III	III	III	III	IX	·	
<i>Galerina pumila</i>	·	4.7	3.5	4.7	8.5	3.5	10.5	·	I	I	I	II	II	VIII	·	
<i>Lycoperdon nigrescens</i>	·	1.7	1.2	·	·	3.3	0.2	·	I	I	·	·	III	I	·	
<i>Macrolepiota procera</i>	·	0.7	1.0	·	1.2	2.8	0.2	·	I	I	·	I	IV	I	·	
<i>Hypholoma fasciculare</i>	·	2.0	·	0.2	0.5	17.7	41.0	·	I	·	I	I	IV	I	·	
<i>Dacrymyces stillatus</i>	·	8.3	·	·	5.0	254.2	861.0	·	I	·	·	I	IV	IX	·	
<i>Mycena metata</i>	·	0.3	·	0.8	0.3	0.2	16.3	·	I	·	I	I	I	VIII	·	
<i>Mycena aetites</i>	·	0.2	·	·	0.3	0.3	0.7	·	I	·	·	I	I	I	·	
<i>Paneolus papilionaceus</i>	·	5.0	0.3	·	·	0.3	·	·	I	I	·	·	I	·	·	
<i>Hygrocybe virginea</i> var. <i>virginea</i>	·	2.3	7.7	0.2	0.5	1.2	·	·	I	I	I	I	II	·	·	
<i>Gymnopus putillus</i>	·	0.2	0.5	0.7	0.5	0.2	·	·	I	I	I	I	I	·	·	
<i>Rhodocybe caelata</i>	·	0.3	1.0	1.5	0.3	0.3	·	·	I	I	I	I	I	·	·	
<i>Lycoperdon perlatum</i>	·	4.2	3.5	7.3	5.8	4.2	·	·	II	I	II	I	IV	·	·	
<i>Flammulaster granulatus</i>	·	2.8	0.3	5.7	0.2	5.3	·	·	I	I	I	I	II	·	·	
<i>Mycena vitilis</i>	·	0.3	0.2	2.7	5.5	1.8	·	·	I	I	I	II	II	·	·	
<i>Mycena stylobates</i>	·	0.3	0.8	0.7	1.2	2.3	·	·	I	I	I	I	II	·	·	
<i>Galerina hypnorum</i>	·	1.5	8.8	4.8	12.2	29.2	·	·	I	I	I	II	VIII	·	·	
<i>Lycoperdon molle</i>	·	5.8	0.7	2.3	·	15.7	·	·	I	I	I	·	VIII	·	·	
<i>Sphaerobolus stellatus</i>	·	0.8	·	0.8	·	25.2	·	·	I	·	I	·	II	·	·	
<i>Clavaria argillacea</i>	·	239.2	193.0	47.3	92.8	·	·	·	VI	V	II	III	·	·	·	
<i>Rhodocybe parilis</i>	·	6.8	8.2	1.5	0.2	·	·	·	II	II	I	I	·	·	·	
<i>Galerina atkinsoniana</i>	·	32.3	7.5	111.5	63.5	·	·	·	III	I	VI	IV	·	·	·	
<i>Hygroaster asterosporus</i>	·	1.3	1.8	13.8	11.8	·	·	·	I	I	IV	III	·	·	·	
<i>Cantharellula umbonata</i>	·	220.0	25.7	55.7	·	·	·	·	IV	I	II	·	·	·	·	
<i>Lycoperdon dermoxanthum</i>	·	4.2	6.7	0.5	·	·	·	·	II	III	I	·	·	·	·	
<i>Entoloma jubatum</i>	·	2.2	4.5	2.0	·	·	·	·	I	II	I	·	·	·	·	
<i>Rhodocybe hirneola</i>	·	0.2	·	2.0	·	·	·	·	I	·	I	·	·	·	·	
<i>Cystoderma jasonis</i>	·	·	0.8	4.7	1.7	·	·	·	·	I	I	I	·	·	·	
var. <i>lilacipes</i>	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	
<i>Clitocybe fragrans</i>	·	·	5.2	4.7	2.5	8.5	·	·	·	I	I	I	VI	·	·	
<i>Hygrocybe conica</i> var. <i>conica</i>	·	·	0.8	1.0	0.8	1.8	·	·	·	I	I	I	II	·	·	
<i>Gymnopus impudicus</i>	·	·	0.7	·	2.2	1.3	·	·	·	I	·	I	II	·	·	
<i>Clitocybe odora</i> var. <i>odora</i>	·	·	1.0	·	·	3.2	·	·	·	I	·	·	I	·	·	

<i>Mycena adonis</i>	.	.	2.8	.	.	.	0.7	.	.	I	.	.	.	I
<i>Cystoderma jasonis</i> var. <i>jasonis</i>	.	.	0.7	.	3.7	.	0.2	.	.	I	.	I	.	I
<i>Collybia cookei</i>	.	.	4.7	.	.	2.5	0.8	.	.	I	.	.	I	I
<i>Collybia tuberosa</i>	.	.	0.3	8.0	1.2	.	14.7	.	.	I	I	I	.	III
<i>Leptosporomyces galzinii</i>	.	.	0.2	1.2	8.0	1.2	22.0	.	.	I	I	II	II	X
<i>Cystodermella granulosa</i>	.	.	.	4.8	0.8	0.2	0.5	.	.	.	I	I	I	II
<i>Strobilurus stephanocystis</i>	.	.	.	2.5	20.3	0.7	0.2	.	.	.	I	III	I	I
<i>Phanerochaete sanguinea</i>	.	.	.	0.5	0.3	0.7	0.7	.	.	.	I	I	I	II
<i>Dacrymyces variisporus</i>	.	.	.	11.8	17.5	41.5	2.0	.	.	.	I	I	V	I
<i>Clitocybe candicans</i>	.	.	.	7.2	0.2	15.2	0.2	.	.	.	II	I	VII	I
<i>Baeospora myosura</i>	.	.	.	0.7	1.5	6.0	3.8	.	.	.	I	I	III	III
<i>Mycena sanguinolenta</i>	.	.	.	0.3	.	11.8	0.3	.	.	.	I	.	IV	I
<i>Mycena vulgaris</i>	.	.	.	12.5	.	.	13.5	.	.	.	I	.	.	III
<i>Clitocybe anisata</i>	.	.	.	.	2.3	15.0	0.7	.	.	.	.	I	V	I
<i>Auriscalpium vulgare</i>	.	.	.	.	0.8	11.0	2.7	.	.	.	.	I	V	IV
<i>Galerina mniophila</i>	.	.	.	.	2.5	.	0.8	.	.	.	.	I	.	II
<i>Coprinopsis friesii</i>	16.5	.	.	.	.	.	.	VI	.	.	.	.	.	.
<i>Parasola plicatilis</i>	5.0	.	.	.	.	.	.	III	.	.	.	.	.	.
<i>Clitocybe brumalis</i>	0.5	.	.	.	.	.	.	I	.	.	.	.	.	.
<i>Conocybe spiculoides</i>	0.5	.	.	.	.	.	.	I	.	.	.	.	.	.
<i>Mycetinis scorodonius</i>	.	45.5	.	.	.	.	.	.	III	.	.	.	.	.
<i>Entoloma rhodopolium</i>	.	0.3	.	.	.	.	.	.	I	.	.	.	.	.
<i>Conocybe rickeniana</i>	.	0.2	.	.	.	.	.	.	I	.	.	.	.	.
<i>Crepidotus variabilis</i>	.	5.8	1.2	.	.	.	.	.	I	I	.	.	.	.
<i>Psathyrella gyroflexa</i>	.	3.2	0.7	.	.	.	.	.	II	I	.	.	.	.
<i>Macrolepiota mastoidea</i>	.	0.3	0.2	.	.	.	.	.	I	I	.	.	.	.
<i>Gymnopus dryophilus</i>	.	0.8	6.0	.	.	.	.	.	I	I	.	.	.	.
<i>Lycoperdon ericaeum</i>	.	.	6.8	.	.	.	.	.	.	I	.	.	.	.
<i>Entoloma conferendum</i>	.	.	2.5	.	.	.	.	.	.	I	.	.	.	.
<i>Lycoperdon marginatum</i>	.	.	2.3	.	.	.	.	.	.	I	.	.	.	.
<i>Lycoperdon norvegicum</i>	.	.	0.7	.	.	.	.	.	.	I	.	.	.	.
<i>Tapinella atrotomentosa</i>	.	.	0.3	.	0.2	.	.	.	.	I	.	I	.	.
<i>Polyporus ciliatus</i>	.	.	0.2	.	0.2	.	.	.	.	I	.	I	.	.
<i>Marasmiellus vaillantii</i>	.	.	.	9.2	.	5.0	.	.	.	.	I	.	II	.
<i>Mycena parabolica</i>	.	.	.	0.3	.	0.8	.	.	.	.	I	.	I	.
<i>Trichaptum fuscoviolaceum</i>	.	.	.	6.5	.	.	.	.	.	.	I	.	.	.
<i>Melanoleuca graminicola</i>	.	.	.	2.5	.	.	.	.	.	.	I	.	.	.
<i>Pholiota gumosa</i>	.	.	.	1.3	.	.	.	.	.	.	I	.	.	.
<i>Calocera furcata</i>	.	.	.	0.8	.	.	.	.	.	.	I	.	.	.
<i>Lycoperdon excipuliforme</i>	.	.	.	0.8	.	.	.	.	.	.	I	.	.	.
<i>Lepiota pseudolilacea</i>	.	.	.	0.3	.	.	.	.	.	.	I	.	.	.
<i>Entoloma nitens</i>	.	.	.	0.2	.	.	.	.	.	.	I	.	.	.
<i>Galerina badipes</i>	.	.	.	0.2	.	.	.	.	.	.	I	.	.	.
<i>Mycena arcangeliana</i>	.	.	.	0.2	.	.	.	.	.	.	I	.	.	.
<i>Psathyrella fibrillosa</i>	.	.	.	0.2	0.2	.	.	.	.	.	I	I	.	.
<i>Ripartites tricholoma</i>	.	.	.	0.8	1.2	.	.	.	.	.	I	I	.	.
<i>Mycena speirea</i>	.	.	.	0.5	15.0	.	.	.	.	.	I	II	.	.
<i>Mycena septentrionalis</i>	.	.	.	.	0.2	.	.	.	.	.	.	I	.	.
<i>Agaricus dulcidulus</i>	.	.	.	.	0.2	0.2	.	.	.	.	.	I	I	.
<i>Clitocybe phyllophila</i>	.	.	.	.	8.3	11.8	.	.	.	.	.	I	IV	.
<i>Hypholoma lateritium</i>	.	.	.	.	0.5	4.0	.	.	.	.	.	I	I	.
<i>Amylostereum laevigatum</i>	.	.	.	.	0.8	10.0	.	.	.	.	.	I	II	.
<i>Cystoderma carcharias</i>	.	.	.	.	0.3	11.8	.	.	.	.	.	I	III	.
<i>Tubaria furfuracea</i>	.	0.5	.	.	.	65.5	.	.	I	.	.	.	VI	.
<i>Gymnopus confluens</i>	.	.	.	2.8	.	145.3	.	.	.	.	I	.	IV	.

<i>Gymnopus aquosus</i>	.	.	.	1.8	.	29.0	.	.	.	.	.	I	.	VI	.
<i>Tremella mesenterica</i>	.	.	.	1.5	.	3.3	.	.	.	.	.	I	.	II	.
<i>Psathyrella candolleana</i>	.	.	.	.	.	16.7	.	.	.	.	.	.	.	V	.
<i>Flammulaster rhombosporus</i>	.	.	.	.	.	25.2	.	.	.	.	.	.	.	IV	.
<i>Mycena stipata</i>	.	.	.	.	.	5.3	.	.	.	.	.	.	.	IV	.
<i>Coprinellus xanthothrix</i>	.	.	.	.	.	1.8	.	.	.	.	.	.	.	IV	.
<i>Peniophora polygonia</i>	.	.	.	.	.	66.2	.	.	.	.	.	.	.	III	.
<i>Exidia plana</i>	.	.	.	.	.	23.7	.	.	.	.	.	.	.	III	.
<i>Clitocybe metachroa</i>	.	.	.	.	.	8.2	.	.	.	.	.	.	.	III	.
<i>Crucibulum laeve</i>	.	.	.	.	.	7.5	.	.	.	.	.	.	.	III	.
<i>Galerina marginata</i>	.	.	.	.	.	6.7	.	.	.	.	.	.	.	III	.
<i>Phaeomarasmium erinaceus</i>	.	.	.	.	.	5.5	.	.	.	.	.	.	.	III	.
<i>Daedaleopsis confragosa</i>	.	.	.	.	.	2.5	.	.	.	.	.	.	.	III	.
<i>Crepidotus lundellii</i>	.	.	.	.	.	85.0	.	.	.	.	.	.	.	II	.
<i>Coprinellus micaceus</i>	.	.	.	.	.	40.0	.	.	.	.	.	.	.	II	.
<i>Stereum hirsutum</i>	.	.	.	.	.	16.2	.	.	.	.	.	.	.	II	.
<i>Hymenochaete tabacina</i>	.	.	.	.	.	11.5	.	.	.	.	.	.	.	II	.
<i>Resupinatus trichotis</i>	.	.	.	.	.	10.5	.	.	.	.	.	.	.	II	.
<i>Phlebia rufa</i>	.	.	.	.	.	9.0	.	.	.	.	.	.	.	II	.
<i>Steccherinum fimbriatum</i>	.	.	.	.	.	8.3	.	.	.	.	.	.	.	II	.
<i>Schizophyllum amplum</i>	.	.	.	.	.	6.8	.	.	.	.	.	.	.	II	.
<i>Rhodocybe fallax</i>	.	.	.	.	.	6.7	.	.	.	.	.	.	.	II	.
<i>Mycena filopes</i>	.	.	.	.	.	6.0	.	.	.	.	.	.	.	II	.
<i>Ceraceomyces crispatus</i>	.	.	.	.	.	2.7	.	.	.	.	.	.	.	II	.
<i>Conocybe subpubescens</i>	.	.	.	.	.	1.5	.	.	.	.	.	.	.	II	.
<i>Mycena rubromarginata</i>	.	.	.	.	.	1.0	.	.	.	.	.	.	.	II	.
<i>Coprinellus disseminatus</i>	.	.	.	.	.	66.0	.	.	.	.	.	.	.	I	.
<i>Schizopora paradoxa</i>	.	.	.	.	.	10.0	.	.	.	.	.	.	.	I	.
<i>Gloeocystidiellum porosum</i>	.	.	.	.	.	8.7	.	.	.	.	.	.	.	I	.
<i>Marasmius epiphyllus</i>	.	.	.	.	.	8.3	.	.	.	.	.	.	.	I	.
<i>Gymnopus acervatus</i>	.	.	.	.	.	5.7	.	.	.	.	.	.	.	I	.
<i>Pleurotus ostreatus</i>	.	.	.	.	.	5.3	.	.	.	.	.	.	.	I	.
<i>Trametes ochracea</i>	.	.	.	.	.	5.3	.	.	.	.	.	.	.	I	.
<i>Hyphoderma praetermissum</i>	.	.	.	.	.	4.8	.	.	.	.	.	.	.	I	.
<i>Vesiculomyces citrinus</i>	.	.	.	.	.	3.3	.	.	.	.	.	.	.	I	.
<i>Byssomerulius corium</i>	.	.	.	.	.	2.7	.	.	.	.	.	.	.	I	.
<i>Oligoporus alni</i>	.	.	.	.	.	2.7	.	.	.	.	.	.	.	I	.
<i>Calocera cornea</i>	.	.	.	.	.	2.3	.	.	.	.	.	.	.	I	.
<i>Tubaria confragosa</i>	.	.	.	.	.	2.2	.	.	.	.	.	.	.	I	.
<i>Fomes fomentarius</i>	.	.	.	.	.	2.0	.	.	.	.	.	.	.	I	.
<i>Psathyrella tephrophylla</i>	.	.	.	.	.	2.0	.	.	.	.	.	.	.	I	.
<i>Marasmius setosus</i>	.	.	.	.	.	1.7	.	.	.	.	.	.	.	I	.
<i>Panus suavissimus</i>	.	.	.	.	.	1.3	.	.	.	.	.	.	.	I	.
<i>Strobilurus tenacellus</i>	.	.	.	.	.	1.3	.	.	.	.	.	.	.	I	.
<i>Clitocybe odora</i> var. <i>alba</i>	.	.	.	.	.	1.2	.	.	.	.	.	.	.	I	.
<i>Clitocybe vermicularis</i>	.	.	.	.	.	1.0	.	.	.	.	.	.	.	I	.
<i>Mycena flavescens</i>	.	.	.	.	.	1.0	.	.	.	.	.	.	.	I	.
<i>Rhodocollybia maculata</i>	.	.	.	.	.	1.0	.	.	.	.	.	.	.	I	.
<i>Lepista nuda</i>	.	.	.	.	.	0.8	.	.	.	.	.	.	.	I	.
<i>Mycena inclinata</i>	.	.	.	.	.	0.7	.	.	.	.	.	.	.	I	.
<i>Phellinus punctatus</i>	.	.	.	.	.	0.7	.	.	.	.	.	.	.	I	.
<i>Athelia epiphylla</i>	.	.	.	.	.	0.5	.	.	.	.	.	.	.	I	.
<i>Calocera viscosa</i>	.	.	.	.	.	0.5	.	.	.	.	.	.	.	I	.
<i>Conocybe ochrostriata</i>	.	.	.	.	.	0.5	.	.	.	.	.	.	.	I	.
<i>Hyphodontia quercina</i>	.	.	.	.	.	0.5	.	.	.	.	.	.	.	I	.



<i>Peniophora cinerea</i>	.	.	.	.	.	0.5	.	.	.	.	.	.	I	.
<i>Clitocybe nebularis</i>	.	.	.	.	.	0.3	.	.	.	.	.	.	I	.
<i>Geastrum quadrifidum</i>	.	.	.	.	.	0.3	.	.	.	.	.	.	I	.
<i>Mycena supina</i>	.	.	.	.	.	0.3	.	.	.	.	.	.	I	.
<i>Chlorophyllum rachodes</i>	.	.	.	.	.	0.2	.	.	.	.	.	.	I	.
<i>Crepidotus cesatii</i>	.	.	.	.	.	0.2	.	.	.	.	.	.	I	.
<i>Hygrocybe laeta</i>	.	.	.	.	.	0.2	.	.	.	.	.	.	I	.
<i>Lepiota magnispora</i>	.	.	.	.	.	0.2	.	.	.	.	.	.	I	.
<i>Resinicium bicolor</i>	.	.	.	.	.	0.2	.	.	.	.	.	.	I	.
<i>Tulasnella pallida</i>	.	.	.	.	.	0.2	.	.	.	.	.	.	I	.
<i>Inonotus rheades</i>	.	.	.	.	.	15.7	0.2	.	.	.	.	.	VII	I
<i>Hyphoderma setigerum</i>	.	.	.	.	.	4.0	0.5	.	.	.	.	.	II	I
<i>Gymnopus peronatus</i>	.	.	.	.	.	0.8	0.5	.	.	.	.	.	I	I
<i>Ceraceomyces sublaevis</i>	.	.	.	.	.	0.5	0.5	.	.	.	.	.	I	I
<i>Tubulicrinis subulatus</i>	.	.	.	.	.	0.2	0.2	.	.	.	.	.	I	I
<i>Gloeoporus taxicola</i>	.	.	.	.	.	0.3	2.2	.	.	.	.	.	I	I
<i>Gymnopus ocior</i>	.	.	.	.	.	0.2	4.3	.	.	.	.	.	I	I
<i>Hyphodontia aspera</i>	.	.	.	.	.	1.0	1.3	.	.	.	.	.	I	II
<i>Coniophora arida</i>	.	.	.	.	.	1.0	2.2	.	.	.	.	.	I	IV
<i>Phlebiella sulphurea</i>	.	.	.	.	.	0.7	3.2	.	.	.	.	.	I	V
<i>Mycena galopus</i>	.	.	1.7	.	.	2.5	23.0	.	.	I	.	.	III	VI
<i>Xeromphalina cornui</i>	.	.	.	.	.	11.2	12.3	.	.	.	.	.	II	VII
<i>Mycena zephirus</i>	.	.	.	.	.	1.2	67.0	.	.	.	.	.	I	VIII
<i>Panellus mitis</i>	.	.	.	.	1.0	.	5.8	.	.	.	.	I	.	I
<i>Hygrophoropsis aurantiaca</i>	.	.	.	.	0.3	.	3.3	.	.	.	.	I	.	IV
<i>Exobasidium vaccini</i>	.	.	.	.	.	.	84.2	.	.	.	.	.	.	X
<i>Hypholoma capnoides</i>	.	.	.	.	.	.	17.5	.	.	.	.	.	.	IV
<i>Botryobasidium vagum</i>	.	.	.	.	.	.	4.0	.	.	.	.	.	.	IV
<i>Mycena purpureofusca</i>	.	.	.	.	.	.	2.3	.	.	.	.	.	.	II
<i>Armillaria ostoyae</i>	.	.	.	.	.	.	1.0	.	.	.	.	.	.	II
<i>Athelia decipiens</i>	.	.	.	.	.	.	0.5	.	.	.	.	.	.	II
<i>Hyphodontia subalutacea</i>	.	.	.	.	.	.	0.5	.	.	.	.	.	.	II
<i>Dacrymyces ovisporus</i>	.	.	.	.	.	.	14.3	.	.	.	.	.	.	I
<i>Pseudomerulius aureus</i>	.	.	.	.	.	.	13.3	.	.	.	.	.	.	I
<i>Dacrymyces macnabbii</i>	.	.	.	.	.	.	8.3	.	.	.	.	.	.	I
<i>Pholiota spumosa</i>	.	.	.	.	.	.	2.3	.	.	.	.	.	.	I
<i>Gymnopilus penetrans</i>	.	.	.	.	.	.	1.7	.	.	.	.	.	.	I
<i>Phlebiella pseudotsugae</i>	.	.	.	.	.	.	1.5	.	.	.	.	.	.	I
<i>Crepidotus applanatus</i>	.	.	.	.	.	.	0.8	.	.	.	.	.	.	I
<i>Gymnopilus bellulus</i>	.	.	.	.	.	.	0.7	.	.	.	.	.	.	I
<i>Galerina calyptrata</i>	.	.	.	.	.	.	0.5	.	.	.	.	.	.	I
<i>Pseudohydnum gelatinosum</i>	.	.	.	.	.	.	0.3	.	.	.	.	.	.	I
<i>Tapinella panuoides</i>	.	.	.	.	.	.	0.3	.	.	.	.	.	.	I
<i>Macrolepiota permixta</i>	.	.	.	.	.	.	0.2	.	.	.	.	.	.	I

Abbreviations and comments: see Tab. 1

clusters 1 and 2, e.g., *Clitocybe diatreta*, *Cystoderma amianthinum*, *Cantharellula umbonata*, *Galerina atkinsoniana*, *Hygroaster asterosporus*, *Psilocybe montana*, *Rickenella fibula*. Some species, like *Clavaria argillacea*, *Rhodocybe parilis*, *Galerina atkinsoniana*, *Hygroaster asterosporus*, *Lycoperdon dermoxanthum*, *Entoloma jubatum*, *Rhodocybe hirneola*, *Cystoderma jasonis* var. *lilacipes* were found only in the communities of psammophilous grassland with juniper and pine (phases 4-6/7) (Tab. 2).

The saprobic species fruiting evenly in a number of succession phases are rather few, e.g., *Mycena aetites*, *Gymnopus putillus*, and *G. impudicus*.

In contrast to the ECM fungi, the saprobic species occurring constantly from the advanced optimal stage, through the juniper-aspen brushwood to the pine forest (terminal stage), are numerous, e.g., *Leptosporomyces galzinii*, *Cystodermella granulosa*, *Strobilurus stephanocystis*, *Phanerochaete sanguinea*, *Dacrymyces variisporus*, *Clitocybe candicans*, *Baeospora myosura* and *Mycena sanguinolenta* (Tab. 2). Also, the group of species exclusive to the terminal stage occurring both in the brushwood and forest community, is relatively rich, e.g., *Hyphoderma setigerum*, *Gymnopus peronatus*, *Hyphodontia aspera*, *Coniophora arida*, *Phlebiella sulphurea*, *Xeromphalina cornui* and *M. zephyrus*. These are the species inhabiting litter and dead wood.

The more developed the vegetation, the higher the number of saprotrophic exclusive species. In the early succession phases their number, abundance and spatial frequency were low (with the exception of *Coprinopsis friesii* and *Parasola plicatilis* in phase 2, and *Mycetinis scorodonius* in phase 4); in the juniper-aspen brushwood and in the pine forest they are frequently much higher.

### 5.2.3. Spatial frequency patterns

Apart from producing different numbers of carpophores in subsequent phases of succession, fungal species show different spatial distribution of their carpophores. For the ECM fungi, usually the increase of carpophore abundance is accompanied by an increase of spatial frequency (Tab. 1); the pattern is less clear in saprotrophs (Tab. 2). Moreover, for many ECM species, after attaining maximum abundance a decrease in carpophore production is not immediately followed by decreasing frequency and the frequency tends to stay high longer. The pattern is distinct in *Amanita muscaria*, *Chalciporus piperatus*, *Tricholoma equestre* and *Boletus edulis* (Tab. 1).

Both in the ECM and non-ECM fungi, species attaining low spatial frequency ( $\leq 30\%$ ; class I-III) prevail. However, proportions of species of low, medium (31-70%; class IV-VII) or high ( $> 70\%$ ; class VIII-X) frequency in subsequent succession phases are different in these groups (Fig. 19A, B). For the ECM species, the relative number of species attaining low spatial frequency decreases in the course of succession process; both the relative numbers of medium and high frequency species tend to increase. For the saprobic species, the proportion of low frequency species is the highest in the optimal stage of succession (phases 5-6/7).

### 5.2.4. Ectomycorrhizal carpophore production

The average annual number of ECM carpophores per 1000 m<sup>2</sup> slightly exceeds a thousand in phases 4 and 5. Then it grows considerably and reaches a maximum level of over 2000 carpophores in phase 6, in a well developed optimal stage of succession with pines under 20 years old, and starts to drop again in the later phase of the optimal stage. In the terminal stage, in transitional juniper-aspen brushwood the number of carpophores decreases down to ca 500 and to 200 in the forest (Fig. 20). As far as the ECM species in the whole successional series are concerned, the most numerous fruit bodies are produced by *Cortinarius*

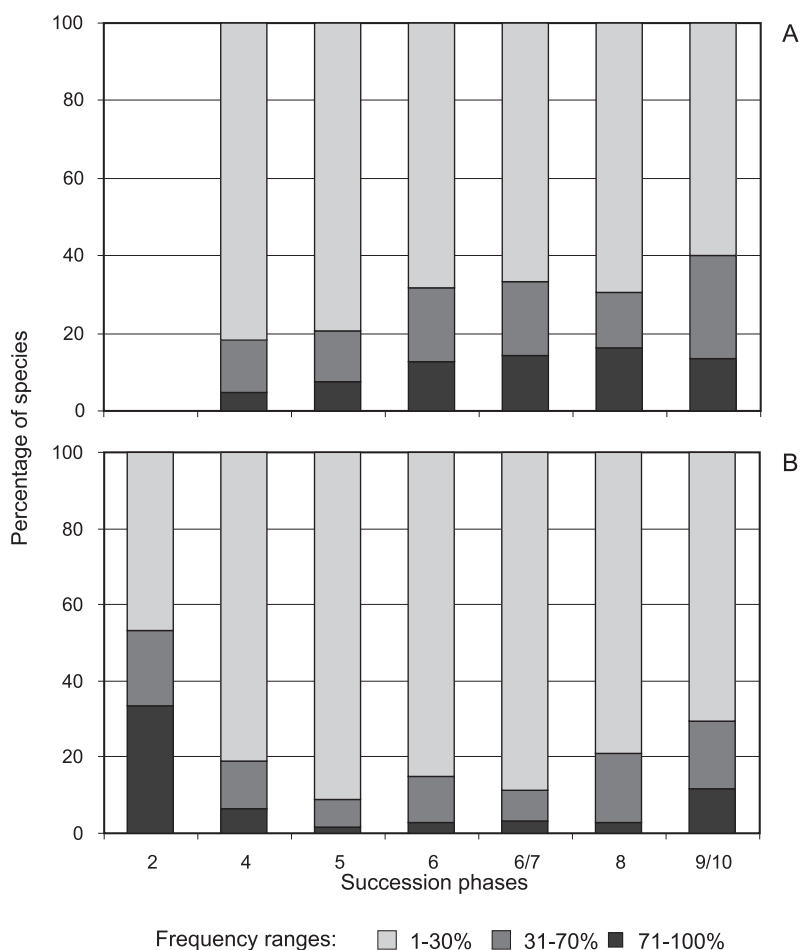


Fig. 19. Distribution of spatial frequency of ECM fungi (A) and non-ECM fungi (B) in consecutive phases of the spontaneous secondary succession on abandoned farmland.

*cinamomeoluteus* and other *Cortinarius* species from the *Cortinarius* I group, *Inocybe lace-  
ra*, *Hygrophorus hypothejus* and *Laccaria proxima*.

The average annual carpophore biomass production (as dry weight) per 1000 m<sup>2</sup> in the ECM fungi is at its lowest in phase 9/10 – 273 g, and at its highest in phase 6 – 1491 g (Fig. 20). This means from 2.7 to 14.9 kg of dry weight per year per ha.

In the season richest in fungal carpophores (out of the three years of observations), in an overall study area of 1.2 ha, 19,917 fruit bodies of ECM fungi were recorded giving a total dry weight of 15.3 kg. According to the calculations by ARNOLDS (1981) this is equivalent to about 153 kg of fresh carpophore biomass. The highest production – ca 21 kg ha<sup>-1</sup>d.w. – was recorded in the plots representing advanced optimal stage of succession, with pines of 8-15 (19) and 12-18(25) years old (phases 6 and 6/7).

The proportion of the total ECM carpophore biomass to their number in consecutive succession phases is much lower in the initial and optimal stages; in the terminal stage (phases 8 and 9/10) it increases considerably. This means that in contrast to open plant

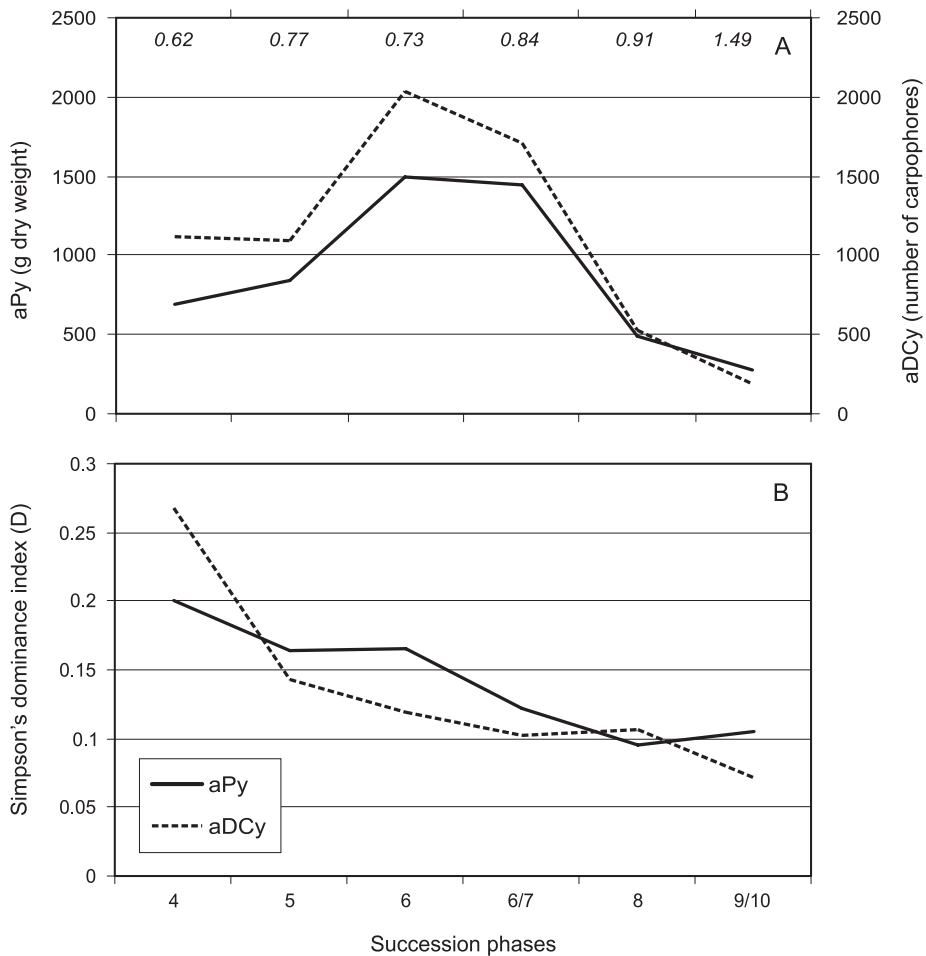


Fig. 20. Average annual carpophore production (aPy) and average annual carpophore density (aDCy) (A), and Simpson's dominance index (D) based on aPy and aDCy (B) of ECM fungi in consecutive phases of the spontaneous secondary succession on abandoned farmland. In (A) figures in italics denote the mean weight of carpophores in the subsequent succession phases.

communities where there is a high production of rather small fungal fruit bodies, in transitional and forest communities species producing bigger carpophores prevail (Fig. 20A). Mean weight of an ECM carpophore (in g d.w.) in the examined succession phases increased from 0.62 in phase 4 to 1.49 in phase 9/10.

The highest carpophore productivity is attained by the genera *Amanita*, *Cortinarius*, *Inocybe*, *Laccaria*, *Russula*, *Suillus* and *Tricholoma*. The proportion of these species in the total annual carpophore production exceeds 90% in the initial communities and still reaches 75% in the pine forest (Fig. 21). The early succession phases (4 and 5) are characterized by high production of the genus *Amanita* (mainly *A. muscaria*, see Tab. 1). The genus *Inocybe* shows clear preference towards initial communities, while *Suillus* and *Laccaria* are typical of open vegetation. In the optimal stage, particularly in its advanced phases (6 and 6/7) the participation of *Tricholoma* becomes significant, in the juniper-aspen brushwood (8)

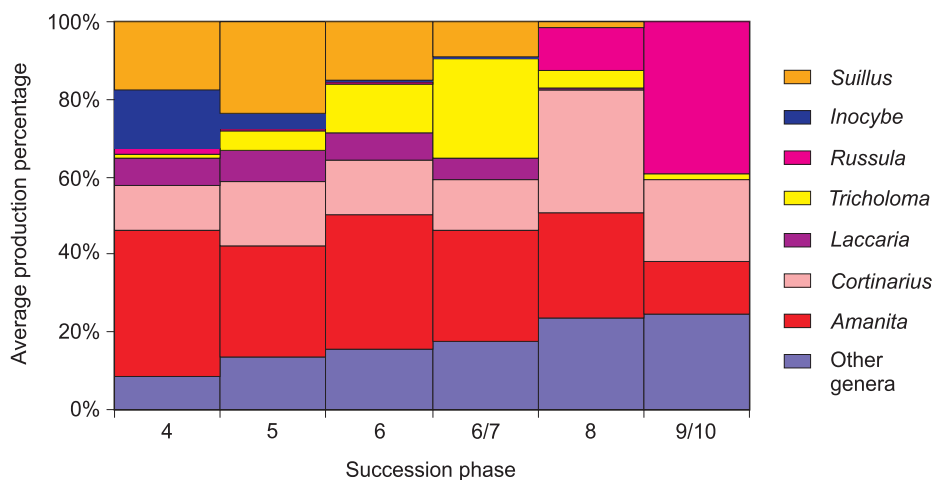


Fig. 21. Percentage of average annual production (aPy) of seven most productive ECM genera in consecutive phases of the spontaneous secondary succession on abandoned farmland.

– *Cortinarius*, and in the pine forest (9/10) – *Russula*. The vegetation development is followed by increasing participation of fungi belonging to other genera. Gradual diversification of the ECM carpophore production along the successional gradient is reflected in the decrease of Simpson's dominance index, both in biomass production and in carpophore numbers (Fig. 20B).

### 5.3. Similarity structure of macrofungi

The similarity structures of the ECM (Fig. 22) and saprobic (Fig. 23) fungal communities occurring in the subsequent succession phases revealed by cluster analysis show high coincidence with the similarity structure of vegetation (Fig. 24) as well as with the model of its succession; the species composition and the carpophore abundance patterns correspond to its subsequent phases (compare Fig. 2). A specificity of the ECM and saprobic fungal succession is also apparent.

The analysis of the species composition and carpophore density of the ECM fungi clearly separates typical forest communities (phase 9/10) from the communities of open and transitional sites (Fig. 22A). Also, it reveals high similarity among ECM communities in the plots representing phases 4 and 5, and among the communities in the plots representing phases 6 and 6/7. It is worth noticing that time (thus the age of symbiotic trees and/or soil development) is not always a decisive factor for similarity structure of fungal assemblages. Moreover, the ECM communities present in the plots occupied by juniper-aspen brushwood seem to be more similar to much earlier communities in phases 4 and 5 (see also Fig. 12). It results from a considerable proportion of species associated with deciduous trees in these plots. However, when assemblages of the ECM fungi growing in the plots of the same phase are combined and treated together, at phase level the analysis shows higher mycological similarity of brushwood and forest (phases 8 and 9/10) (Fig. 22B).



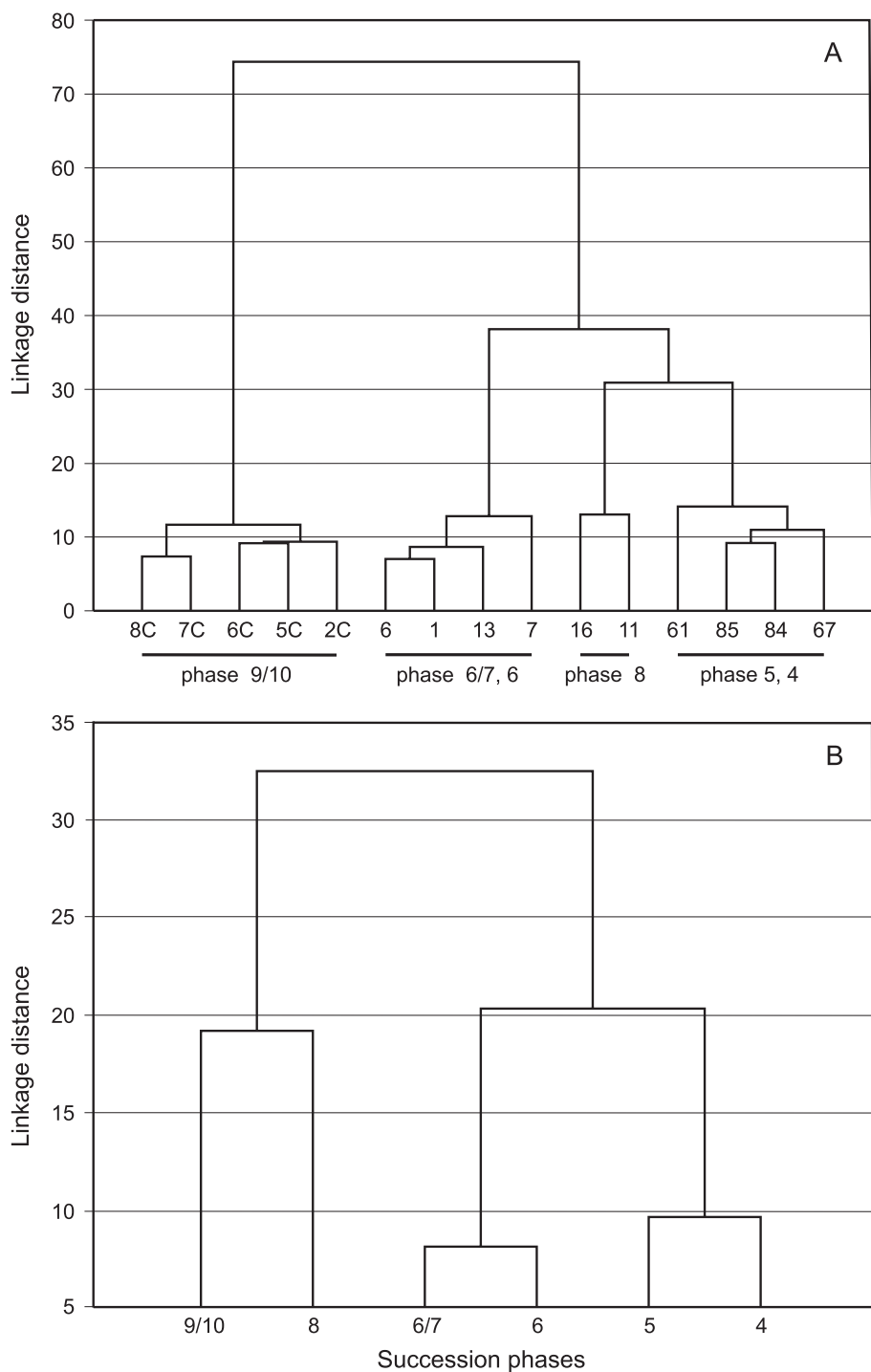


Fig. 22. Similarity structure of the examined plots (A) and succession phases (B) based on ECM species composition and annual density of carpophores.

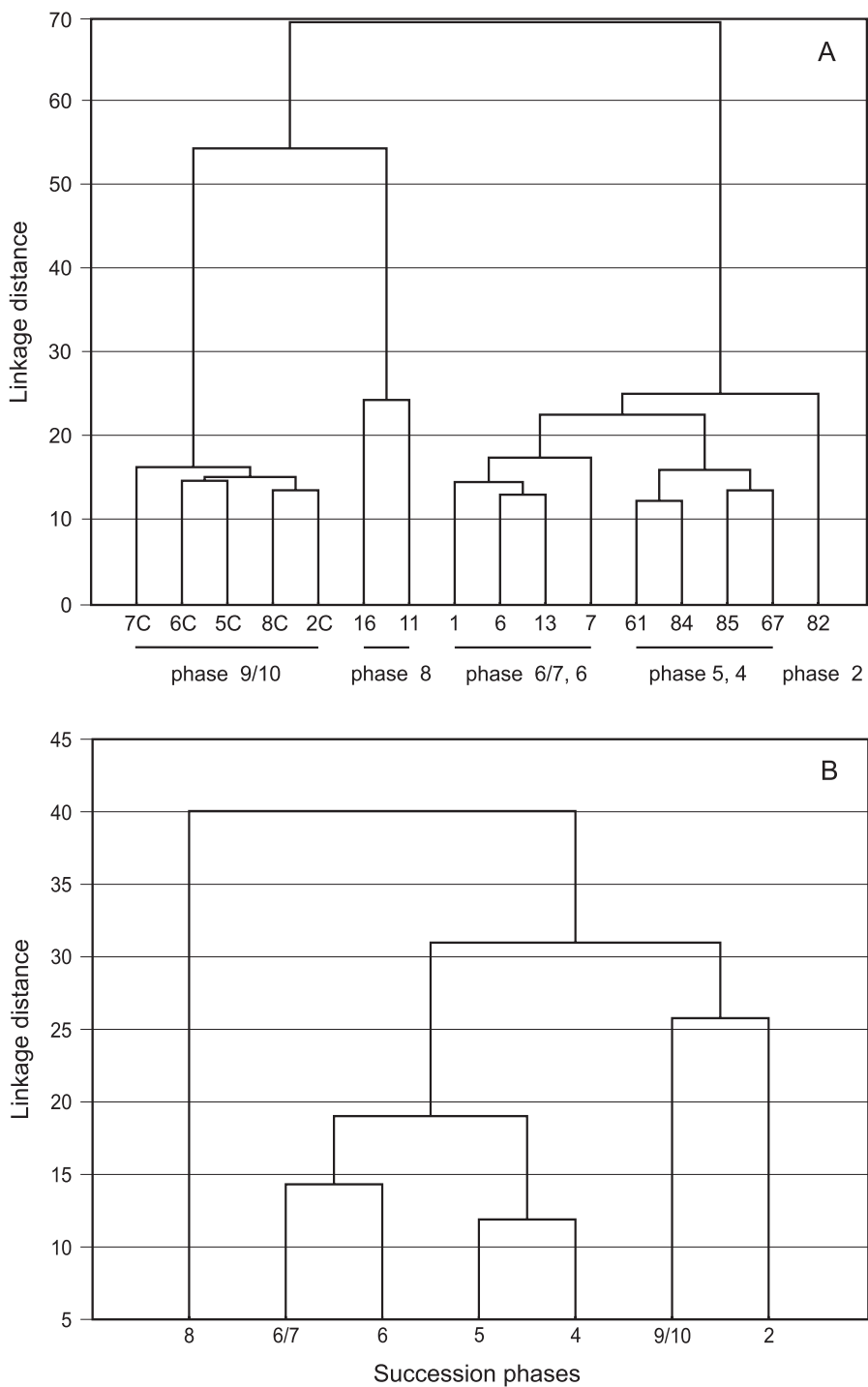


Fig. 23. Similarity structure of the examined plots (A) and succession phases (B) based on non-ECM species composition and annual density of carpophores.

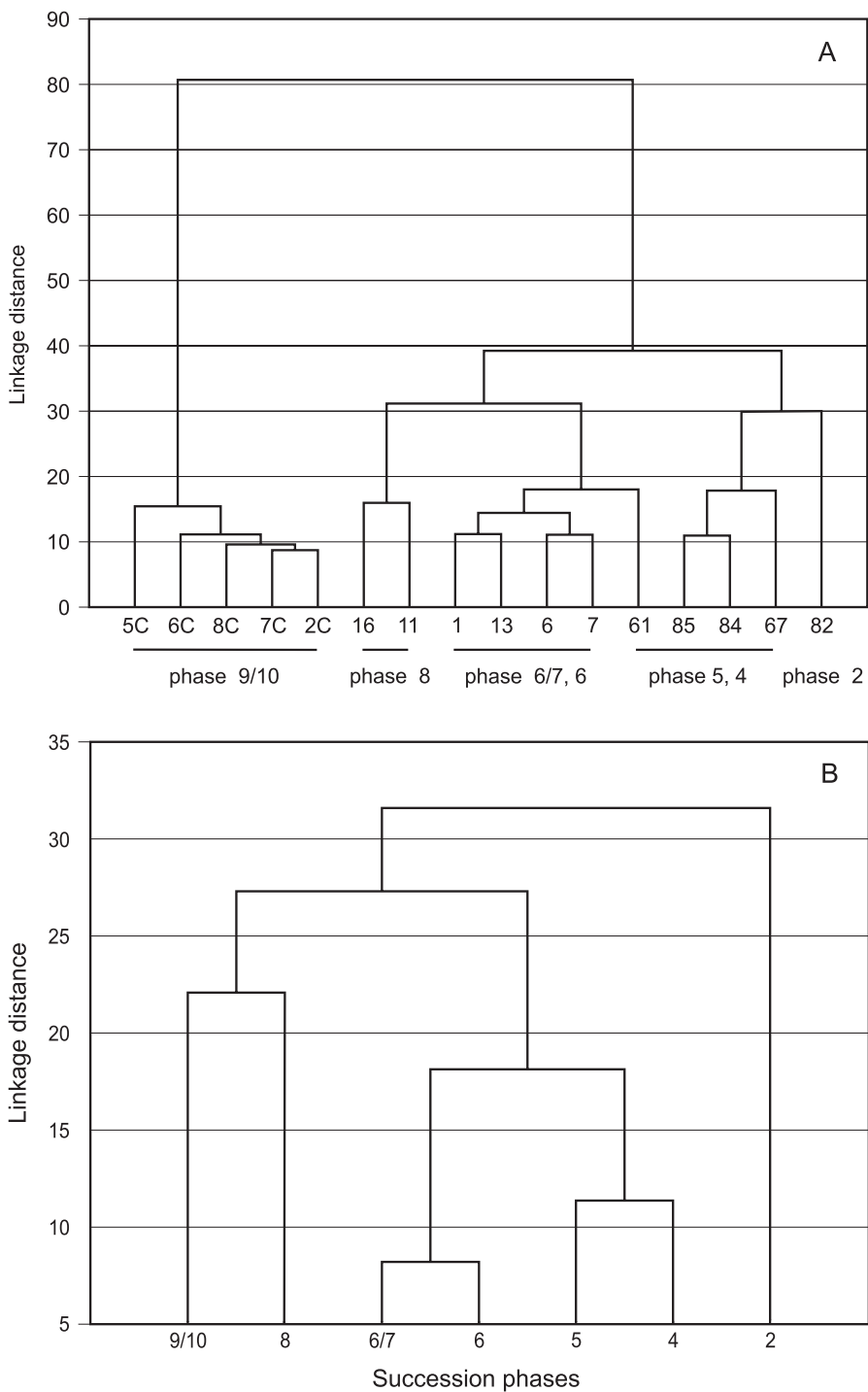


Fig. 24. Similarity structure of the examined plots (A) and succession phases (B) based on vegetation composition and cover.

The analysis of similarity structure of the non-ECM species reveals almost the same pattern as in the ECM fungi in initial and optimal stage of succession (Fig. 23A), but in contrast, the communities of saprotrophic fungi in the plots with juniper-aspen brushwood reveal higher similarity to the communities growing in forest. The analysis clearly separates the assemblages of non-ECM fungi of phase 2, phase 8 and phase 9/10; this probably reflects totally different kinds and quality of available substrates.

#### **5.4. Quantitative changes in ectomycorrhizal roots along successional gradient**

Examination of the ECM roots in surface soil samples revealed some differences in their vertical distribution. In early succession phases, the roots grow mainly in the upper 3-5 cm of mineral soil increasing considerably its stabilization. Both in initial and optimal stage, in sites of litter accumulation the ECM roots occur mainly in the interface between mineral soil and organic matter; the extramatrical mycelium frequently overgrows and binds the litter pieces together. In phase 6/7, where the percentage of dead roots of pine and juniper in the surface soil is the highest, the living roots are tightly interwoven with masses of the dead ones and the hyphae emanating from the ECM mantles make it impossible to separate them. The roots and mycelium are also present in great abundance in places of humus horizon formation. In the juniper-aspen brushwood, the main body of ECM roots overgrows the humus layer and the adjacent part of litter. In the pine forest, the roots penetrate mainly the Of and Oh horizons.

Although the qualitative structure of the ectomycorrhizae is not a subject of the present study, the observations of ECM roots during the quantitative examination revealed the increasing mycorrhizal diversity along the successional gradient – the greatest number of morphotypes was found in the juniper-aspen brushwood. There is also an increase in the systems' ramification; simple or dichotomous morphotypes prevail in the earlier phases while in the latter phases also multi-ramified and coralloid systems are frequent.

The length of living fine roots and the number of ECM tips in soil samples also change in the subsequent succession phases, showing high within-phase variation (Tab. 3).

The highest length of the living ECM roots in the surface soil was found in the juniper-aspen brushwood (phase 8), with an average of 54.7 and 35.7 cm per 100 cm<sup>3</sup> of soil in two consecutive seasons. The number of mycorrhizal tips is also the highest – ca 305 per 100 cm<sup>3</sup> of soil. The figures are significantly higher than in the other succession phases.

Despite considerable differences in the ECM tip number and the living root length among particular samples, plots and phases, the ratio of these two parameters, i.e. ramification index, was relatively stable in roots sampled in the same year independently of the succession phase or species composition of a tree-stand. The average index was 4.1-5.6 and 7.1-9.2 in two consecutive seasons (Tab. 3). The correlation between the root length and the number of mycorrhizal tips is high – 0.90 and 0.92, respectively (Spearman's rank correlation coefficient;  $P < 0.0001$ ).

The number of well developed (WD) mycorrhizas varied considerably among different samples, succession phases and between sampling seasons; a relatively low proportion of WD mycorrhizas in phase 6/7 may be a result of lower soil moisture in these plots (Appendix 1, Fig. A1.1).

Table 3

Length of living tree roots and total number of ECM root tips per 100 cm<sup>3</sup>, number of ECM tips per cm root length, and number of well developed (NWD) ECM tips relative to the total number of ECM tips in the soil samples ( $\pm$ S.D.)

Year	Succession stage	Succession phase	Living root length (cm/100 cm <sup>3</sup> )	Number of ECM tips (n/100 cm <sup>3</sup> )	Number of ECM tips / cm root	NWD ECM tips (%)
1997	initial	2	7.9 $\pm$ 1.6	34.3 $\pm$ 29.9	4.1 $\pm$ 2.9	46.49 $\pm$ 21.34
		4	21.6 $\pm$ 14.2	107.2 $\pm$ 75.5	5.0 $\pm$ 1.6	77.58 $\pm$ 14.53
	optimal	5	18.5 $\pm$ 15.4	90.8 $\pm$ 73.9	5.0 $\pm$ 1.6	49.46 $\pm$ 23.61
		6	29.4 $\pm$ 17.7	138.8 $\pm$ 75.4	4.9 $\pm$ 1.1	62.13 $\pm$ 19.20
		6/7	16.1 $\pm$ 7.8	81.1 $\pm$ 53.5	5.0 $\pm$ 1.6	54.38 $\pm$ 17.18
	terminal	8	54.7 $\pm$ 16.6*	304.5 $\pm$ 94.6*	5.6 $\pm$ 1.1	77.75 $\pm$ 15.36
		9/10	19.2 $\pm$ 5.7	94.6 $\pm$ 48.6	5.1 $\pm$ 2.3	73.16 $\pm$ 11.83
1998	initial	2	5.4 $\pm$ 2.1	45.8 $\pm$ 8.3	8.9 $\pm$ 1.9	87.78 $\pm$ 2.48
		4	12.0 $\pm$ 6.8	106.7 $\pm$ 60.6	9.2 $\pm$ 3.7	66.83 $\pm$ 19.36
	optimal	5	8.2 $\pm$ 5.0	72.6 $\pm$ 49.6	8.6 $\pm$ 1.3	77.02 $\pm$ 14.55
		6	11.3 $\pm$ 9.6	91.5 $\pm$ 67.5	9.0 $\pm$ 2.1	60.78 $\pm$ 20.83
		6/7	12.7 $\pm$ 6.0	90.0 $\pm$ 39.9	7.6 $\pm$ 2.1	39.33 $\pm$ 18.50
	terminal	8	35.7 $\pm$ 16.8*	304.8 $\pm$ 147.5*	8.9 $\pm$ 1.8	76.27 $\pm$ 10.09
		9/10	8.4 $\pm$ 2.7	58.1 $\pm$ 18.5	7.1 $\pm$ 1.7	75.79 $\pm$ 14.88

\* Figures statistically different from all the other figures of the same year within a column (Tukey HSD test,  $p < 0.01$ )

The average frequency of ECM tips on roots exceeded 95% in all the plots, including phase 2, where no ECM sporocarps were found yet; the roots of young birches growing as one solitary group in this plot were colonized by the ECM fungi in 98.6%.

## 6. DISCUSSION

**Sporocarp sampling.** Observations of sporocarps, their taxonomical determination, and estimation of frequency and abundance remain one of the main methods in the studies of species richness and diversity, distribution and ecology of macromycetes. The presence of sporocarps in the field is always evidence of the presence of vegetative mycelium of the species and this is the fundamental merit of this approach. However, the lack of generative structures or the lack of records of such structures does not necessarily mean that the mycelium of a certain species is not present at a site (ARNOLDS 1992b). Sporocarps are frequently ephemeral and impermanent, their production can vary considerably depending on unpredictable weather fluctuations (amount of rainfall, temperature, humidity), on seasons and a few years' cycles. For these reasons, studies based on sporocarp occurrence

should span long-term observations, 3 to 8 years, depending on the size of monitored area and the frequency of sporocarp recording (ARNOLDS 1981, 1992a, b; BARKMAN 1987; VOGT *et al.* 1992; WATLING 1995; FRIEDRICH 2008).

The present study is based on a 3 year period of sporocarp observations and includes 17 recording visits for each plot (except for plot 82, phase 2, where 13 visits were performed in two seasons). Most of the visits were carried out in the periods of intensive sporocarp production (autumn) so that the maximum diversity and numbers of fruit bodies could be registered. All three seasons were relatively favourable for carpophore production. The permanent plots comprised altogether an area of 13,000 m<sup>2</sup>; to increase the exactitude of observations and to examine the sporocarp spatial distribution, the plots were divided into subplots (25-100 m<sup>2</sup>) searched one by one, individually. One cycle of visits to all the plots took 2-6 days. This field work resulted in a 55% increase in the ECM species number during the second observation season but only 3% increase during the third season; for the non-ECM species these figures were 27% and 6%, respectively. Continuing the sporocarp surveys in the plots for further years could blur the differences between the studied succession phases as the successional changes, especially in the vegetation of phases 4-6, were very fast.

### **6.1. Macrofungi in the scheme of spontaneous secondary succession on the abandoned fields near Białowieża**

**Fungi in the initial stage of vegetation succession.** First macrofungi probably appear very early in the succession process, inhabiting dead plant remnants and places colonized by mosses on fresh fallow (phase 1) soon after the field abandonment. In the herb-grassland community (phase 2) 15 saprotrophic basidiomycete species were found, among which *Crinipellis scabellia*, *Marasmius oreades*, *Mycena flavoalba*, *M. olivaceomarginata* and *M. leptcephala* prevail. These are the species characterised by broad ecological range, known from various plant communities rich in herbaceous and grass debris (ARNOLDS 1992c). High abundance and frequency of their occurrence results from considerable homogeneity of the available substrate, both in terms of its composition (abundant necromass of plant shoots and leaves decaying on compact turf of dead and living roots) and spatial distribution (high and even field layer cover). The presence of bryophilous fungi, like *Galerina* spp., *Cystoderma amianthinum* and *Rickenella fibula*, which accompany mosses entering the fallow, is also noticeable.

An abundant presence of agricultural weeds or even legumes can considerably slow down or even cease the succession process on post-arable fields (FALIŃSKI 1986b; VAN DER PUTTEN *et al.* 2000; HEDLUND & GORMSEN 2002). This phenomenon can be seen in plot 82, where compact field layer dominated with *Poa pratensis*, *Elymus repens*, *Stellaria graminea*, *Vicia* spp. and *Agrostis* spp., along with a thick turf of roots and herbaceous debris, has prevented juniper and tree colonization for more than 30 years after the field abandonment – the plot still represents phase 2 of the vegetation succession.

The appearance of pioneer tree seedlings, mainly of pine, which according to Faliński's model takes place in phase 2 (usually ca 5 years after field abandonment, see Fig. 2) is crucial both for the further development of plant communities towards forest and for fungal assemblages. The ECM inoculum potential of the soil in this phase, necessary for successful tree colonization (TRAPPE & LUOMA 1992; BOERNER *et al.* 1996) seems high despite the short



time after the cultivation cessation. Close vicinity to older communities with trees of various age and especially to the great species reservoir of the Białowieża old-growth forest probably contribute to the fast restoration of the ECM spore bank in the soil similar to the constant and considerable inflow of seeds from the forest plants (SYMONIDES 1986; FALIŃSKI *et al.* 1993). The position of a site in the close vicinity of a forest or the presence of single trees or groups of trees are among the most important primary and secondary factors affecting the successional trajectory of meadow-type vegetation (FALIŃSKA 2003).

Pine seedlings usually emerge in places where the plant or lichen cover has been disrupted by animals (e.g., *Myrmecoleon*, wild boars) or among naturally fragmented tufts of mosses (FALIŃSKI 1986b). The rate of successional processes seems to be particularly high in sites of poor psammophilous grassland character that neighbour field strips much more advanced in terms of vegetation development; i.e. juniper-aspen brushwood or nearly closed-canopy pine stand. Tree seedlings emerging close to older trees benefit via increased mycorrhizal colonization (DICKIE & REICH 2005; DICKIE *et al.* 2007). The facilitation is even more efficient if seedlings appear near the compact forest tree-line and are incorporated into the common mycorrhizal network of hyphae growing from the roots of mature trees (contagion model), which potentially reduces the costs of establishing mycorrhizal infection (SIMARD & DURALL 2004; THIET & BOERNER 2007; TESTE *et al.* 2009). In contrast, seedlings emerging within a dense community of tall grass and herbs are subject to strong herbaceous competition regardless of distance from trees (DICKIE *et al.* 2007). This might also be one of the reasons why the succession process and trees colonization in plot 82 (phase 2) have been arrested.

In phase 4, young, mostly 6-8 year old trees are accompanied by over 40 symbiotic ECM fungal species producing epigeous sporocarps. A majority of them are more or less host-specific to pine, even in places where birch, aspen and willow are also abundant. This might confirm a 'coniferous' direction of the vegetation succession similar to the presence of pioneer lichen species – *Cladonia cornuta* and *C. gracilis* (FALIŃSKI *et al.* 1993). The highest carpophore abundance and very high spatial frequency is attained by *Inocybe lacera*, *Suillus luteus* and *Amanita muscaria*. Numerous carpophores occur of *Cortinarius*, especially *Cortinarius cinnamomeoluteus*, and also *Laccaria proxima*, *Hygrophorus hypothejus* and *Chalciporus piperatus*. These species form a group, which later constantly accompanies plant communities in optimal stage of succession, producing very abundant and frequent sporocarps. They are still present in the juniper-aspen brushwood but their abundance and frequency there is low. The highest production in terms of carpophore biomass in the initial stage is attained by the genera *Amanita* (mainly *A. muscaria*), *Suillus* and *Inocybe*.

The ECM species comprise nearly half of the fungi occurring in phase 4. A relatively low diversification of available substrates and a high cover of field and moss layers do not favour saprotrophic fungi, which are not as numerous as in the later phases. The species growing on soil and among mosses prevail: *Psilocybe montana*, *Clitocybe diatreta*, *Clavaria argillacea*, *Cystoderma amianthinum*, *Rickenella fibula*. The species inhabiting litter, which is not abundant yet, are rather few; similarly, the species inhabiting wood occur sporadically. Most species found in the initial stage also occur in the optimal stage of the vegetation succession.

*Inocybe lacera* and *I. asterospora*, the species attaining maximum abundance and frequency in phase 4, as well as *Cantharellula umbonata*, *Mycetinis scorodoni* and the species of *Entoloma*, which were found exclusively in both plots of phase 4, can be regarded as distinctive species for this phase.

**Fungi in the optimal stage of vegetation succession.** As far as fungi are concerned, the borderline between the communities of the last phase of the initial stage and the early phase of the optimal stage is not distinct. The number of both ECM and saprotrophic fungal species increases. Besides the fungi known from phase 4, a few ECM species, such as *Laccaria bicolor* and species of *Tricholoma*, considerably increase their presence. A group of new species attaining the highest abundance and, in most cases, high frequency in the optimal stage appear, for instance *Cortinarius mucosus*, *C. parevernus* and *Lactarius rufus*. The proportion of species occurring with high spatial frequency increases and the dominance index decreases. For phase 6, when pines are 8-15(19) years old, considerably higher abundance, production and spatial frequency of the following combination of species is distinctive: *Hygrophorus hypotheus*, *Amanita muscaria*, *Chalciporus piperatus*, *Laccaria proxima* and *L. bicolor*. On the other hand, the presence of a few species exclusive for the optimal stage, such as *Inocybe curvipes*, *Cortinarius anthracinus*, *C. mucosus*, *C. colus*, *Tricholoma saponaceum* and *Sacrodon squamosus*, is also characteristic. *Lactarius rufus* and *Xerocomus badius* are the only two ECM species which start fruiting in the optimal stage of the vegetation development and still occur in the pine forest; the abundance of the first one is the highest around canopy closure, the latter one clearly prefers mature forest conditions. The number of carpophores and the productivity of the ECM fungi reach their maximum, the species richness is the highest at plot level (38 species on 1000 m<sup>2</sup> in both plots). This coincides with reaching maturity by the open community of psammophilous grassland with rich moss-lichen layer, junipers and pines (*Spergulo-Corynephorum cladinetosum mitis*) (FALIŃSKI *et al.* 1993). It is interesting that in phase 6 the number of lichen species and the cover of lichens is also the highest in the chronosequence studied (FALIŃSKI *et al.* 1993).

Later, when pines are 12-18 to 25 years old, the crowns of trees growing in large groups tend to meet and form a canopy in places. Together with big juniper shrubs they intercept considerable amount of rainfall, while additionally very dense roots drain the soil. As a result xeric conditions are created (FALIŃSKI *et al.* 1993). Shade increases as does the production and accumulation of litter. At the same time the number of ECM carpophores, especially small ones, decreases considerably; the productivity and species richness is also slightly lower. Very abundant occurrence of different species of *Tricholoma* is distinctive for this phase of vegetation development, marking a specific period of canopy closure, intensified litter accumulation but still the lack of a true humus layer and very low C and N content in the mineral superficial soil (Appendix 1).

In the group of saprotrophic fungi, the species which appeared in phase 4 still prevail in the optimal stage. Most of them attain maximum carpophore abundance. Early in this stage, numerous species of "gasteromycetes" occur, such as *Lycoperdon molle*, *L. nigrescens*, *L. perlatum*, *L. ericaeum*, *L. marginatum*, *L. norvegicum* (the last three species were found exclusively in phase 5), *Lycoperdon dermoxanthum* (maximum abundance and frequency in phase 5) and *Scleroderma fuscum*. In close vicinity of the plot representing this phase *Lycoperdon utriforme* was also found. High species richness of this group of fungi seems to be distinctive for phase 5 of the vegetation development.

Bryophilous species are very abundant and frequent in the optimal stage, e.g., species of *Galerina* and *Cystoderma amianthinum* in large tufts of *Polytrichum commune* and *P. juniperinum*, as well as *Psilocybe montana* and *Rickenella fibula* in the synusia of *Polytrichum piliferum*. Some other species typical of open areas of psammophilous grasslands, heaths and juniper thickets (BREITENBACH & KRÄNZLIN 1986-2000; BAS *et al.* 1988-1995) grow

abundantly on soil, mainly in moss-lichen patches, e.g., *Hygroaster asterosporus*, *Entoloma jubatum*, *Hygrocybe virginea* and a few species of *Rhodocybe*.

As the amount and diversity of organic debris increases, species inhabiting litter (mainly pine and juniper needles) and wood become more and more numerous. Great numbers of them also occur in the later succession phases, in the juniper-aspen brushwood and pine forest.

**Fungi in the terminal stage of vegetation succession – juniper-aspen brushwood.** The brushwood community is characterised by the highest species richness and the highest overall diversity of macromycetes.

Changes of the species composition of the ECM fungi in comparison with the previous stages are relatively minor, while quantitative changes are distinct. Numerous species, which occurred earlier, cease production of their carpophores and occur with low abundance and frequency. In the case of some species, for instance *Amanita muscaria*, *Chalciporus piperatus*, *Tricholoma equestre* and *Boletus edulis*, initially the number of carpophores is strongly limited and later also the spatial frequency decreases. Such a phenomenon was also observed in the brushwood communities for psammophilous mosses (FALIŃSKI *et al.* 1993).

Species that appear in the juniper-aspen brushwood for the first time and are still present in the pine forest are few. These are *Cantharellus cibarius*, *Russula paludosa* and *Amanita citrina*.

Many ECM species are exclusive for the juniper-aspen brushwood, some of which achieve high carpophore abundance. As compared with the earlier phases, the number of species host-specific to deciduous trees increases considerably. The following species are distinctive for this phase: *Cortinarius trivialis*, *C. alboviolaceus*, *Russula clavipes*, *Lactarius uvidus* and *L. torminosus*.

The species richness of ECM fungi at phase level is the highest. In contrast, the number of ECM carpophores, as well as their biomass production, is much lower than in the optimal stage, approaching the values observed in the forest. The genus *Cortinarius* prevails in terms of the number of species, carpophore abundance and production.

The species composition of saprotrophic fungi changes considerably; almost 50% of species found in the juniper-aspen brushwood are exclusive for this community. This may be a consequence of the dominance of deciduous trees and of exceptional resource heterogeneity and niche diversification resulting from high cover and density of undergrowth shrubs and trees, humid microclimate and plenty of variable substrates originating from both conifer and broadleaved trees. The majority of saprotrophic species (ca 80%) inhabit litter and wood. The proportion of bryophilous species is relatively low (ca 3%) although they are still present in scattered moss patches.

A shift in the dominance structure of saprotrophic species can be observed. The species attaining high abundance and low spatial frequency as well as the species attaining low abundance and high spatial frequency are much more numerous. In the earlier succession phases the species which were frequent were usually also abundant.

A few species known to occur predominantly in humid, even wet and boggy places (also in the flood plain and riparian vegetation) were found, such as *Russula paludosa*, *Gymnopus aquosus*, *Lentinus suavissimus*, *Marasmius epiphyllus*, *Mycena supina*, *Phaeomarasmium erinaceus*. They correspond to a group of higo- and hydrophilous mosses distinctive for the juniper-aspen brushwood (FALIŃSKI *et al.* 1993), from among which *Amblystegium serpens*, *Brachythecium salebrosum* and *Brachytheciastrum velutinum* were recorded in the

plots examined. The presence of such species results from specific microclimatic conditions: high cover of shrubs and trees, deep shade and relatively high soil humidity.

**Fungi in the terminal stage of vegetation succession – nearly mature fresh pine forest.** In comparison with the earlier succession phases, the number of species decreases, especially of the ECM fungi – their number is lower than in phases 4 and 5. Much lower is also the abundance of carpophores and their biomass production. In contrast to the optimal stage, there are no ECM species producing hundreds of fruit bodies, their spatial distribution is scattered but relatively even.

After mainly quantitative changes in the ECM sporocarp community in the earlier phases, the change associated with the development of pine forest is much more of qualitative character. About 50% of the ECM species found there are new for the chronosequence studied. These are species occurring mainly in pine and mixed forests (with pine as co-dominant tree), e.g., *Amanita porphyria*, *Cortinarius caperatus*, *C. semisanguineus*, *Suillus variegatus*, *Lactarius tabidus*, *Tylopilus felleus*. The most distinctive is an occurrence of numerous species of *Russula*, attaining the highest biomass production of any genus. The presence of *Russula emetica*, *R. paludosa* and *R. vinosa*, species occurring usually in humid or mossy forests, indicates a mesophilous character of the habitat.

In contrast, 73% of saprotrophic species present in the pine forest were growing also in open or pre-forest vegetation phases. Some of them occur less abundantly, such as *Mycena flavoalba*, *Clitocybe diatreta*, *Psilocybe montana*, *Macrolepiota procera*, which are species preferring open communities and likely to disappear in mature forest. On the other hand, the abundances of many species inhabiting litter and pine wood, e.g., *Gymnopus androsaceus*, *Mycena galopus*, *M. zephirus*, *Xeromphalina cornui* and *Dacrymyces stillatus*, *Leptosporomyces galzinii*, *Coniophora arida* are the highest in the pine forest. However, some of these fungi, similarly to some ECM species and a considerable group of typically forest mosses (FALIŃSKI *et al.* 1993), appear as early as in phases 4 and 5 of vegetation development, usually around pines. The majority of the saprotrophic species occurring exclusively in the forest are wood-inhabiting taxa.

## 6.2. Patterns of fungal succession in post-agricultural land

Among the main causes and mechanisms of succession applying both to vegetation and fungi are an availability of space and availability of species of differential performance. Thus the whole process strongly depends on arrival and establishment of species (PICKETT *et al.* 1987; FRANKLAND 1992). The cessation of agricultural practices and field abandonment, especially when the land is liberated from any direct human impact and permanently left aside, creates a space for undisturbed, spontaneous secondary succession. Spores present in large quantities in the air are the most common and readily available supply of different fungal species. The arrival of spores on a particular substrate is usually a random process, although it depends on, e.g., distance from a source, topography, season, fungal life cycles and dispersal mechanisms (FRANKLAND 1992). Potential fungal colonists are affected by factors associated with the resource or substrate quality, microclimate and other organisms (BODDY 1992).

Forming symbiosis with proper ECM fungi is necessary for effective land colonization by trees in boreo-temperate bioms. To make it possible, the presence of inoculum in form of

spores, sclerocia or mycelium is crucial (TRAPPE & LUOMA 1992; MOLINA *et al.* 1992). Arable soil usually lacks ECM inoculum potential and it may only appear some years after the cultivation cessation (BOERNER *et al.* 1996). However, millions of spores present in the air as well as spores left by numerous mycofagous animals can quickly rebuild a dormant spore bank, especially in areas where forests are in close vicinity (ALLEN M.F. 1987; CÁZARES & TRAPPE 1994; JUMPPONEN *et al.* 2002; ASHKANNEJHAD & HORTON 2006; MENKIS *et al.* 2007).

### Species richness, diversity and carpophore production

The proximity of forests, among them the Białowieża old-growth forest, the presence of more advanced vegetation communities on neighbouring field strips which were abandoned earlier as well as management history (extensive agriculture and low mineral fertilizing) resulted in relatively fast natural tree regeneration. Pines, aspens and birches were usually establishing as early as 2 to 5 years after field abandonment (FALIŃSKI 1986a, b; FALIŃSKI *et al.* 1993). Non-disturbed vegetation development also resulted in accumulation and increasing diversification of substrates for saprobic macrofungi. In the open communities where pines were 6-8 to 12 years old, carpophores of as much as 92 macrofungal species were recorded, among them 44 ECM species. After this initial shift in species number, the ECM species richness stabilized and showed only minor increases and fluctuations in open and pre-forest vegetation. The highest overall species richness (183 species) was observed in the juniper-aspen brushwood, mainly because of an exceptionally high number of saprotrophs (134 species). Finally, the number of species in the pine forest decreased considerably both in mycorrhizal and saprotrophic fungi to 30 and 67 taxa, respectively.

According to FALIŃSKI (1986b), natural and undisturbed development of Central European fresh pine forest *Peucedano-Pinetum* lasts about 140 years before achieving maturity, stable species composition and proper vertical structure. The data on the species richness of fungal community in old-growth forest of this type are inconsistent.

In a 27 ha patch of the old-growth forest of the same type but of primeval origin growing in the Białowieża National Park, a total of 288 macrobasidiomycete species were found (non-regular, 4 years examination within research plot V-100) including 100 ECM species, 93 litter-inhabiting species and 93 species growing on wood (BUJAKIEWICZ 1997; LISIEWSKA 1997a, b, c, d; SKIRGIELLO 1997a, b, c, d, e, f). The number of epigeous species (ascomycetes and basidiomycetes, including ECM, soil and litter-inhabiting fungi) found per hectare fluctuated between 10 and 43 (26 on average) (ŻARNOWIEC *et al.* 1996) and the average number of epixylous species was 13 (CHLEBICKI *et al.* 1996). This would mean that in the old-growth pine forest the number of fruiting fungal species found per hectare is even lower than in 0.2 ha in the nearly mature 100-year-old first-generation spontaneous stand growing on former arable land (69 epigeous species including 30 ECM and 39 soil- and litter-inhabiting saprotrophic fungi, and 28 species growing on wood).

On the other hand, in the same type of old-growth forest, approximately in the same part of the Białowieża National Park, but in the years 1955-1956, NESPIAK (1959) found 87 and 80 species of basidiomycete cup-fungi alone in two plots, 100 m<sup>2</sup> each, and in their vicinity (altogether 124 species). Such species richness would be much higher than the richness found in the present study. Difference in species numbers between the two studies in the Białowieża old-growth may be attributed to the high year-to-year variation of carpophore production caused by different weather conditions, to spatial heterogeneity of the old-growth and of fungal assemblages and, to some extent, to different accuracy in examining small (100 m<sup>2</sup>) and large (1 ha) plots, on regular and non-regular basis. Considering the



negative processes affecting fungi in the European forests (ARNOLDS 1991; SCHMITT 1991; TERMORSHUIZEN & SCHAFFERS 1991; FELLNER 1993) it cannot be excluded that this difference reflects a general trend of loss of biodiversity, especially severe in the case of ECM fungi in old stands.

The number of carpophores produced both by the ECM and saprotrophic fungi as well as the ECM carpophore biomass production increased along the old-field successional gradient till achieving its maximum in the optimal stage of vegetation development, in the complex of psammophilous grassland with juniper shrubs and pines mainly 8-15 to 20 years old. The figures decreased slightly around the canopy closure (pines 12-18 to 25 years old) and then decreased conspicuously in the terminal stage, especially in the pine forest.

Similar observations concerning changes of the species richness and carpophore production in the *Pinus sylvestris* stands in Finland were made by HINTIKKA (1988). He compared the species numbers and abundances of the ECM and saprotrophic fungi (excluding species growing on wood) in plantations representing four age classes and found the lowest average number of species and carpophores in the youngest, partly open, sapling stands of pines 5-15 years old. Both species richness and carpophore abundance increased rapidly in the pole stands (20-30 years old), were slightly lower in thinning stands (30-50 years old) and decreased in the stands over 70 years old. TERMORSHUIZEN (1991) found much higher ECM species richness and carpophore abundance in young stands (4-13 years old) than in older stands (50-80 years old) of *Pinus sylvestris* in the Netherlands.

An initial increase of species number and abundance of ECM fungi up to its maximum in the middle-aged stands and a decrease of these parameters in older stands were found also for other tree species. RICEK (1981) observed a gradual increase of the number of species to 29 in the age class 15-20 years (canopy closure) and up to 48 in the age class 20-25 years in the stands of *Picea abies*. In older stands (40-45 years old), the species richness decreased again. For *Pseudotsuga menziesii*, JANSEN (1991) found the highest numbers of species and sporocarps (16 and 5500, respectively) in young stands 8-18 years old (right before canopy closure) and their gradual decrease in older stands, 20-36 and 41-54 years old, down to 8 species and 280 sporocarps in the latter case. In the Norway spruce plantation on formerly agricultural land, GÁPER & LIZOŇ (1995) observed an increase in the number of ECM fungi and their abundance from 7 species and 341 sporocarps in the first three years after tree planting to 11 species and 3967 sporocarps in years 7-10 and a further increase to 19 species but a decrease to less than 400 carpophores in years 24-27 (older stands were not included).

Successional changes in the carpophore production mean both changes in carpophore numbers and in their standing biomass. In the present study, the average annual carpophore biomass production by the ECM fungi increased from zero in phase 2 (lack of ECM sporocarps) to 6.9 kg ha<sup>-1</sup> of dry weight (d.w.) in phase 4 and to 14.9 kg ha<sup>-1</sup> d.w. in phase 6 (pines 8-15 to 20 years old), then it slightly decreased around the canopy closure and dropped considerably to 4.9 kg ha<sup>-1</sup> d.w. in the juniper-aspen brushwood and to 2.7 kg ha<sup>-1</sup> d.w. in the pine forest with ca 100 years old stand. The figures are underestimated as even if the observations are carried out in the peak season at weekly intervals 10-30% of the total sporocarp production could be missed (RICHARDSON 1970).

In the earlier succession phases, as much as nearly 70% of the ECM annual carpophore biomass was produced by 3 species (*Amanita muscaria*, *Suillus luteus*, *Cortinarius cinnamomeoluteus*). In the later phases, the dominance index decreased considerably and in the pine forest, 3 most productive species (*Russula badia*, *Cortinarius caperatus*, *Paxillus*



*involutus*) accounted for about 44% of the ECM annual biomass. Similar results were obtained by DAHLBERG *et al.* (1997) in the Nowary spruce forest of the same age: the 3 most productive species (all of them were *Cortinarius* spp.) accounted for about 44% of the average annual carpophore biomass production by all ECM fungi.

GARDES & BRUNS (1996) found that the ECM species producing abundant carpophores do not always produce abundant mycorrhizae and that the most abundant mycorrhizae may belong to the species producing sparse carpophores or even not producing them at all. Thus the ECM species differ in the way they allocate their biomass between these structures. According to DAHLBERG *et al.* (1997), in the Norway spruce forest in Sweden, the proportion of carpophore biomass in the overall species biomass (carpophore and mycorrhizae) can vary considerably among species, for instance for *Russula decolorans* it equals 0.6-0.9% while for *Cortinarius brunneus* – 10-16%. Moreover, the annual carpophore biomass production (8.8 kg ha<sup>-1</sup> d.w.) comprised less than 5% of the biomass production of mycorrhizae (250-400 kg ha<sup>-1</sup> d.w.). In turn, WALLANDER *et al.* (2001) revealed that 80% of the total biomass of the ECM fungi may belong to the extramatrical mycelium growing in soil; it can amount to 700 kg ha<sup>-1</sup> in the pine stand. The following question arises: if the sporocarp biomass production of the ECM fungi assemblage associated with 10-20 years old pines is over five times greater than that of the assemblage in the pine forest with 100 year old trees, does it mean that the below-ground biomass of mycorrhizae and extramatrical mycelium follows the same pattern or that the allocation of the biomass into sporocarps in the younger, successional stands is bigger than in mature stands. It should be kept in mind that, as in the present study, at least some of the ECM species occur in both age ranges.

The maximum number of ECM carpophores and their biomass was found in phase 6, when pines were about 10-20 years old and their cover reached 50% (compare Appendix 2, Tab. A2.1). The culmination of the ECM production in oligotrophic pine plantations may occur a bit later, under 20-30 years old trees (HINTIKKA (1988). The period of stand development, when trees gain their size and approach the state of canopy closure, seems to have significant consequences for the development of fungal assemblages (DIGHTON & MASON 1985; DIGHTON *et al.* 1986; JANSEN 1991; VISSER 1995). It is thought also to be a period of the highest photosynthesis activity of trees, the highest growth rate and the highest needle area (compare HINTIKKA 1988; TROFYMOW *et al.* 2003; SIMARD *et al.* 2004). In the stands of *Pseudotsuga menziesii*, the biomass of epigeous ECM carpophores was ca 6 times greater in the age classes 30-35 and 45-50 years than in the class >400 years (SMITH *et al.* 2002); the first 50 years is the time of the fastest growth of the Douglas fir. There is a high correlation between the total ECM carpophore biomass and the rate of photosynthesis and annual tree growth rate in individual host-fungus associations (HÖGBERG *et al.* 2001; NARA *et al.* 2003a). On the other hand, nutrient supply (mainly N and P) of trees, which depends largely on the symbiotic ECM fungi, may significantly affect their photosynthesis level; thus this bi-directional relationship and feedback can be of great importance for both partners (NARA *et al.* 2003a).

However, an increase of the species richness along with increasing tree age and developing vegetation was also reported, for example in the subalpine Norway spruce forest (but with an almost doubled sporocarp production in the younger stand) (SENN-IRLET & BIERI 1999), in road-side verges (KEIZER & ARNOLDS 1994) and stands of *Tsuga heterophylla*-*Pinus contorta* (KRANABETTER *et al.* 2005). In the latter case, however, the difference between the species numbers in the stands 20 and 60 years old was much bigger than in the stands over 100 and over 200 years old, in which the number of species was growing

only very slowly. The number of species can also stabilise after canopy closure as in stands of *Pseudotsuga menziesii* (with peak productivity in middle-aged stands; SMITH *et al.* 2002). In a chronosequence of *Pinus banksiana* regenerating after wildfire, VISSER (1995) found a rapid increase in the species richness of the ECM fungi (based both on sporocarp and ectomycorrhizae occurrences) between 6 and 41 year old stands which then stabilised in 65 and 122 year old stands.

Comparisons of the species numbers and abundances among different studies should be made with caution as they refer to different plot sizes, different tree species and different stand histories. However, both the age of trees and the stand structure with a kind of turning point around canopy closure seem to be the key factors for the successional changes of the ECM fungi.

Another important determinant of the fungal species richness is the species richness of trees. According to SCHMIT *et al.* (2005), it accounts for nearly 30% of the variation in the species richness of macrofungi as a group and between 9% and 19% of the variation in the species richness for subgroups of ectomycorrhizal fungi, wood-inhabiting fungi and fungi growing on soil or litter. Communities with high tree species richness are correspondingly characterized by high macrofungal species richness. In the present study, such an effect was also observed. In plots 84 (phase 4) and 85 (phase 5), where the proportions of different deciduous trees were higher than in plots 67 and 61 (phases 4 and 5, respectively), the species richness of fungi was also higher. The richest fungal species composition was found in the juniper-aspen brushwood, where the proportion of deciduous trees and their species number was the highest. An exceptional increase in species numbers was observed for saprotrophic species but the effect was also clear in the group of mycorrhizal ones. At least ten species of ECM fungi would probably not have occurred at this stage of vegetation development if the succession had run without the presence of deciduous trees, especially aspen, in the stand (successional trajectory also present in the Jelonka reserve; the vegetation in plots 13 (phase 6), 1 and 6 (phase 6/7) is likely to develop in such a way; compare Appendix 2, Tab. A2.1). In such cases, the species richness of the ECM fungi in the early terminal stage of vegetation development is probably lower than in the juniper-aspen brushwood (49 species) but higher than in the mature pine forest (30 species). It may also be slightly lower than at the time of canopy closure (42 species).

An increase or decrease in the species richness does not necessarily mean the same changes in the species diversity, as the diversity is affected by the species evenness. In the subsequent succession phases examined in the present study, gradual increase in species diversity (Shannon's diversity index) was observed with higher rates of change in the earlier communities (till phase 6) and slowing down thereafter. The ECM fungal diversity was the highest in the terminal pine forest with pines ca 100 year old (compare Fig. 10), despite the number of species being lower than in the earlier succession phases. This results from the increasing evenness in the carpophore abundances in the ECM fungi assemblages. However, at plot level (1000 m<sup>2</sup>), the highest diversity in the pre-forest vegetation was attained in phase 6/7 with pines 12-18 up to 25 years old, which is the time of attaining distinct canopy closure. Then it slightly decreased but in the mature pine forest it increased again. This finding supports the results of a study in the *Pinus contorta* plantation claiming the highest ECM diversity in successional systems occurring around canopy closure (DIGHTON & MASON 1985; DIGHTON *et al.* 1986). It should be kept in mind that the number of species included in this study was limited to the prevailing ones (Shannon's diversity index is relatively sensitive to the number of 'occasional' species), the oldest stands were 27 years old

and that the examined plots were relatively small. It seems that the ECM species diversity is area-dependent and may achieve its maximum in stands in the later stages of succession, after the canopy closure, if the area examined is large enough to cover the potential species richness and the large-scale habitat homogeneity. According to SÅSTAD (1995), in a homogeneous vegetation of the *Vaccinio-Pinetum* type pine forest the spatial patterns of fungal distribution did not deviate from a random distribution. TWIEG *et al.* (2007), who examined the diversity of the ECM fungi (based on ECM root tips) in four age classes of *Pseudotsuga menziesii* stands, found the greatest increase in the 5- to 26-yr-old age class, a period corresponding with tree canopy closure, and only a minor increase thereafter, with similar diversities in 65- and 100-yr-old stands. However, the authors claim that especially in older stands sampling was inadequate to account for all species present.

The saprotrophic fungi achieve maximum diversity in the juniper-aspen brushwood. This is in accordance with the observations in deciduous and coniferous forests in Canada (VILLENEUVE *et al.* 1989), where the diversity index was much higher in the former communities mainly because of the higher number of saprotrophic species, whereas the numbers of the ECM species were comparable. A significant correlation between the number of litter colonizing saprotrophs and tree species richness was also found in conifer plantations by FERRIS *et al.* (2000). In phase 8, about 80% of species were litter- and wood inhabiting fungi. Such a high number of saprobes probably resulted from the high cover and diversification of the litter layer containing decaying remnants of herbaceous plants, fallen leaves and needles, twigs and bark pieces from different species of trees and shrubs, cones, etc, as well as abundant dead branches lying on the ground or still attached to trees, dying junipers, and a few although still solitary logs. Large amounts of fine woody debris, typical of dense, close stands of relatively young trees, can considerably increase the number of species present at a site (NORDÉN *et al.* 2004). In the juniper-aspen brushwood, true litter horizon was observed for the first time in the chronosequence studied, at some places it was even differentiated into Oi and Of horizons. Moreover, for the first time thin humous horizon of initial mull type or initial moder developed (Appendix 1). Accumulating litter and its composition, depth, moisture, presence of different stages of decomposition and time-related changes in chemical composition positively affect the richness of basidiomycetes (DIX & FRANKLAND 1987; LISIEWSKA 1992; OSONO & TAKEDA 2001; OSONO *et al.* 2006). Soil characteristics, particularly of the humus layer, are also important variables for saprotrophic fungi community structure (TYLER 1989; RÜLING & TYLER 1990). According to ARNOLDS (1981, 1992c) a remarkable relation exists between organic matter content in the soil and sporocarp productivity: in the communities outside forest, the highest productivity was observed in grasslands on soils with rather low organic matter content.

The effect of woody debris on the species richness and abundance of saprotrophic fungi was also conspicuous in the pine forest. The number of wood inhabiting species was the highest there; they produced numerous carpophores and comprised a large group of exclusive species. Large numbers of them inhabited coarse woody debris in different stage of decay: numerous pine branches and larger pieces of wood lying on the ground, logs and stumps (Appendix 5). The amounts of coarse woody debris as well as the size of logs and wood diameter are positively related to fungal species richness and basidiocarp production of wood-inhabiting fungi and can strongly affect their community structure (BADER *et al.* 1995; RENVALL 1995; HØILAND & BENDIKSEN 1996; LINDBLAD 1998; FERRIS *et al.* 2000; NORDÉN *et al.* 2004; URCELAY & ROBLEDO 2009). Species richness is also reported to increase with progressive wood decay (RENVALL 1995; HØILAND & BENDIKSEN 1996). Relatively high

soil moisture as well as high cover of field and moss layers (Appendix 1 and 2) seem to create favourable microclimatic conditions, which are among the most important abiotic variables affecting this group of fungi (BODDY 1992).

### Spatial distribution

Differences in the carpophore spatial frequencies of individual species among the plots representing subsequent succession phases are an indication of their preferences and specificity towards the stage of vegetation development. The presence of carpophores approximately reflects the distribution of the species below-ground (JANSEN 1984; GUIDOT *et al.* 2001; ZHOU *et al.* 2001; HIROSE *et al.* 2004), although their first appearance indicates rather the beginning of fruiting and not necessarily the species establishment.

The spatial distribution of the ECM species along the old-field chronosequence studied suggests that succession of these fungi starts simultaneously with trees colonisation and can be impressively fast. Moreover, as the observed secondary succession of vegetation was entirely natural and spontaneous, all the ECM fungi were indigenous; possible interference of nursery species is excluded. The initial species composition, resulting most probably from the establishment of numerous populations of the same and different species from spore deposits over a relatively short time (not more than 6-8 years after the arrival of first cohort of trees, but probably earlier), tends to persist for several to tens of years (Fig. 25, compare Figs 14, 15). This corresponds to the 'initial floristics' model proposed by EGLER (1954) for secondary successional community development. In this model, all species present in the entire chronosequence may be present soon after disturbance, but different groups of species prevail over time. First populations of nearly 50% of the ECM species present in this series were able to establish during the first years of succession, many of them long before attaining their maximum abundance and spatial frequency. An establishment of some species from the vegetative mycelia spreading from the neighbouring, more developed vegetation patches was highly probable, such as: *Amanita citrina*, *Russula galochroa*, *R. clavipes*, *R. emetica*.

As the youngest stands examined in this study comprised 6-8(12) year old trees and they hosted not less than 33 ( $\pm 7$ ) ECM species/1000 m<sup>2</sup>, a strict group of the earliest colonizers forming symbioses with tree seedlings cannot be distinguished based on the spatial frequency alone. A majority of these species can be regarded as early arriving multistage fungi. Among them the group of species present along the whole chronosequence (phases 4-9/10) can be distinguished as well as groups of species which withdraw after canopy closure or before the mature forest formation.

The species richness is built primarily by an accumulation of species, with a distinct although not numerous group of species specific to the optimal stage of vegetation development, followed by partial but considerable replacement best seen between brushwood and forest phase of the terminal succession stage. As a result many species typical of open vegetation as well as species continuing their fruiting in brushwood community disappeared and a group of species specific to old-growth forests gradually established. Similar observations were made, e.g., by VISSER (1995), BRADBURY *et al.* (1998), KRANABETTER *et al.* (2005) and TWIEG *et al.* (2007), who also found the accumulation of species being the major way of building the ECM species composition in successional stands but only a few species being replaced. According to KRANABETTER *et al.* (2005), in the western hemlock – lodgepole pine old-growth forests (with trees over 200 years old) the frequency of sporocarps was evenly divided between species established in early- and mid-seral stages of development

(40% each), followed by species associated with late-seral stands (20%). However, in the present study the replacement of species was much more intensive: species not occurring earlier comprised 50% of the ECM assemblage in the nearly mature pine forest.

Apart from the “core” species associated with pine, a group of “peripheral” species host-specific to deciduous trees were found throughout the studied series, mainly in the juniper-aspen brushwood. The “peripheral” species associated with small populations of paper birch or spruce independently of the main stand development were also found by KRANABETTER *et al.* (2005).

The species persistence and shifts in their spatial distribution over time seem to depend on their life histories (i.e. dispersal, abilities to colonize trees of different age, physiological and nutritional demands, competitive abilities, etc); at least some species show clear preferences for younger trees/earlier developmental phases, such as *Inocybe lacera* and *Suillus luteus*, for the communities around canopy closure, such as *Tricholoma albobrunneum* and *T. portentosum*, or to mature trees/stands, such as *Paxillus involutus*, *Russula badia* and other species.

For the saprotrophic species similar patterns seem to apply (Fig. 26). The majority of species that were not sporadic were multistage fungi, showing preference for the substrates that were continuously available in the whole chronosequence or its part: herbaceous plant debris, moss-lichen synusia, moss tufts, litter and woody debris. The replacement of species was clearly connected with changing resource types and quality; the main changes in species composition took place after the post-fallow herb-grass community transformation into the psammophilous grassland with juniper and pine, after the canopy closure and in the mature forest community. Similar to the “peripheral” ECM fungi, a group of species inhabiting deciduous litter and wood can be distinguished. The changes resemble to some extent those in successional roadside verges (KEIZER 1993b) with species specific to open grassland (*Mycena flavoalba* group), bryophilous species and a distinct group of wood inhabiting fungi.

Fungal dynamics – establishing new species, changes in sporocarp production and their spatial distribution, is a result of population processes both at inter- and intra-specific level. As summarised by AMEND *et al.* (2009), there are species forming large genets (i.e. continuous mycelial individuals of the same genotype) likely propagating vegetatively, such as *Suillus bovinus* (DAHLBERG & STENLID 1990, 1994) and *Xerocomus* spp. (FIORE-DONNO & MARTIN 2001), as well as species growing as much smaller genets which rather propagate via spores, as it was found, e.g., in *Hebeloma cylindrosporum* (GUIDOT *et al.* 2002), *Laccaria amethystina* (FIORE-DONNO & MARTIN 2001) and *Lactarius xanthogalactus*, *Russula cremoricolor* and *Amanita francheti* (REDECKER *et al.* 2001). Both small and large genets can be formed by *Russula brevipes* (BERGEMANN & MILLER 2002). In some species the size of genets positively correlates with forest development and tree age, but the number of genets decreases (DAHLBERG 1997). For *Suillus bovinus*, estimated numbers of 700-5700 genets ha<sup>-1</sup> in younger (15-20 years old) stands and 30-120 ha<sup>-1</sup> in older (70-140 years old) stands were found (DAHLBERG & STENLID 1994). The maximum size of genets was 1.7-5.3 m in the younger forest and to 17.5 m in the older ones. At the same time the production of sporocarps per unit area decreased. In contrast, *Laccaria amethystina* formed small genets also in mature forests (FIORE-DONNO & MARTIN 2001). Limited carpophore numbers and still large sizes of genets in older forests suggest their fragmentation into several ramets (i.e. spatially discontinuous mycelial units of the same genotype) (DAHLBERG & STENLID 1995; SAWYER *et al.* 2001). Such a process could be an explanation why in some species, for

Stages and phases of succession	Initial stage		Optimal stage			Terminal stage	
	2	4	5	6	6/7	8	9/10
Range of pine age	-	6-8(12)	8-12(16)	8-15(19)	12-18(25)	20-30(35)	~ 100
<i>Laccaria proxima</i>							
<i>Cortinarius</i> spp. I							
<i>Thelephora terrestris</i>							
<i>Rhodocollybia butyracea</i> f. <i>asema</i>							
<i>Cortinarius</i> spp. II							
<i>Russula fragilis</i>							
<i>Cortinarius</i> spp. III							
<i>Paxillus involutus</i>							
<i>Amanita rubescens</i>							
<i>Inocybe lacera</i>							
<i>Suillus luteus</i>							
<i>Cortinarius cinnamomeoluteus</i>							
<i>Hygrophorus hypothejus</i>							
<i>Amanita muscaria</i>							
<i>Chalciporus piperatus</i>							
<i>Tricholoma albobrunneum</i>							
<i>Tricholoma portentosum</i>							
<i>Tricholoma equestre</i>							
<i>Leccinum aurantiacum</i>							
<i>Boletus edulis</i>							
<i>Clitopilus prunulus</i>							
<i>Amanita pantherina</i>							
<i>Hebeloma</i> sp. 1							
<i>Inocybe calida</i>							
<i>Inocybe praetervisa</i>							
<i>Laccaria bicolor</i>							
<i>Cortinarius purpurascens</i>							
<i>Inocybe assimilata</i>							
<i>Inocybe asterospora</i>							
<i>Inocybe curvipes</i>							



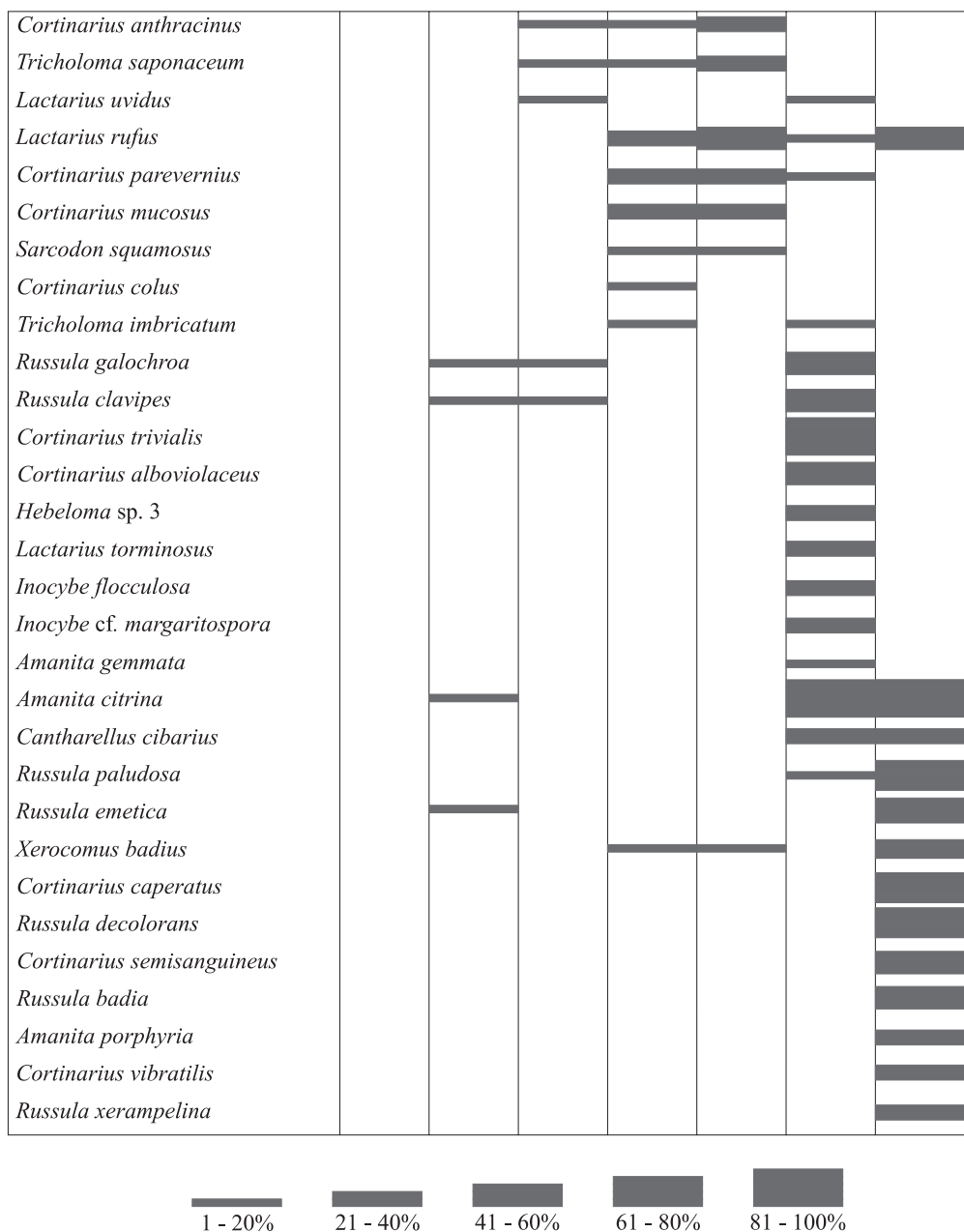


Fig. 25. Spatial frequency of ECM fungi in consecutive phases of the spontaneous secondary succession on abandoned farmland; sporadic species occurring with very low abundance or only in one plot are excluded.

Stages and phases of succession	Initial stage		Optimal stage			Terminal stage	
	2	4	5	6	6/7	8	9/10
Range of pine age	-	6-8(12)	8-12(16)	8-15(19)	12-18(25)	20-30(35)	~ 100
<i>Coprinopsis friesii</i>							
<i>Crinipellis scabella</i>							
<i>Rickenella fibula</i>							
<i>Marasmius oreades</i>							
<i>Mycena olivaceomarginata</i>							
<i>Mycena leptocephala</i>							
<i>Mycena flavoalba</i>							
<i>Galerina</i> spp.							
<i>Clitocybe diatreta</i>							
<i>Cystoderma amianthinum</i>							
<i>Mycena pura</i>							
<i>Psilocybe montana</i>							
<i>Entoloma</i> spp.							
<i>Ampulloclitocybe clavipes</i>							
<i>Clitocybe vibecina</i>							
<i>Gymnopus androsaceus</i>							
<i>Collybia cirrhata</i>							
<i>Galerina pumila</i>							
<i>Dacrymyces stillatus</i>							
<i>Mycena metata</i>							
<i>Lycoperdon nigrescens</i>							
<i>Macrolepiota procera</i>							
<i>Hypholoma fasciculare</i>							
<i>Galerina hypnorum</i>							
<i>Lycoperdon molle</i>							
<i>Lycoperdon perlatum</i>							
<i>Clavaria argillacea</i>							
<i>Galerina atkinsoniana</i>							
<i>Hygroaster asterosporus</i>							
<i>Cantharellula umbonata</i>							

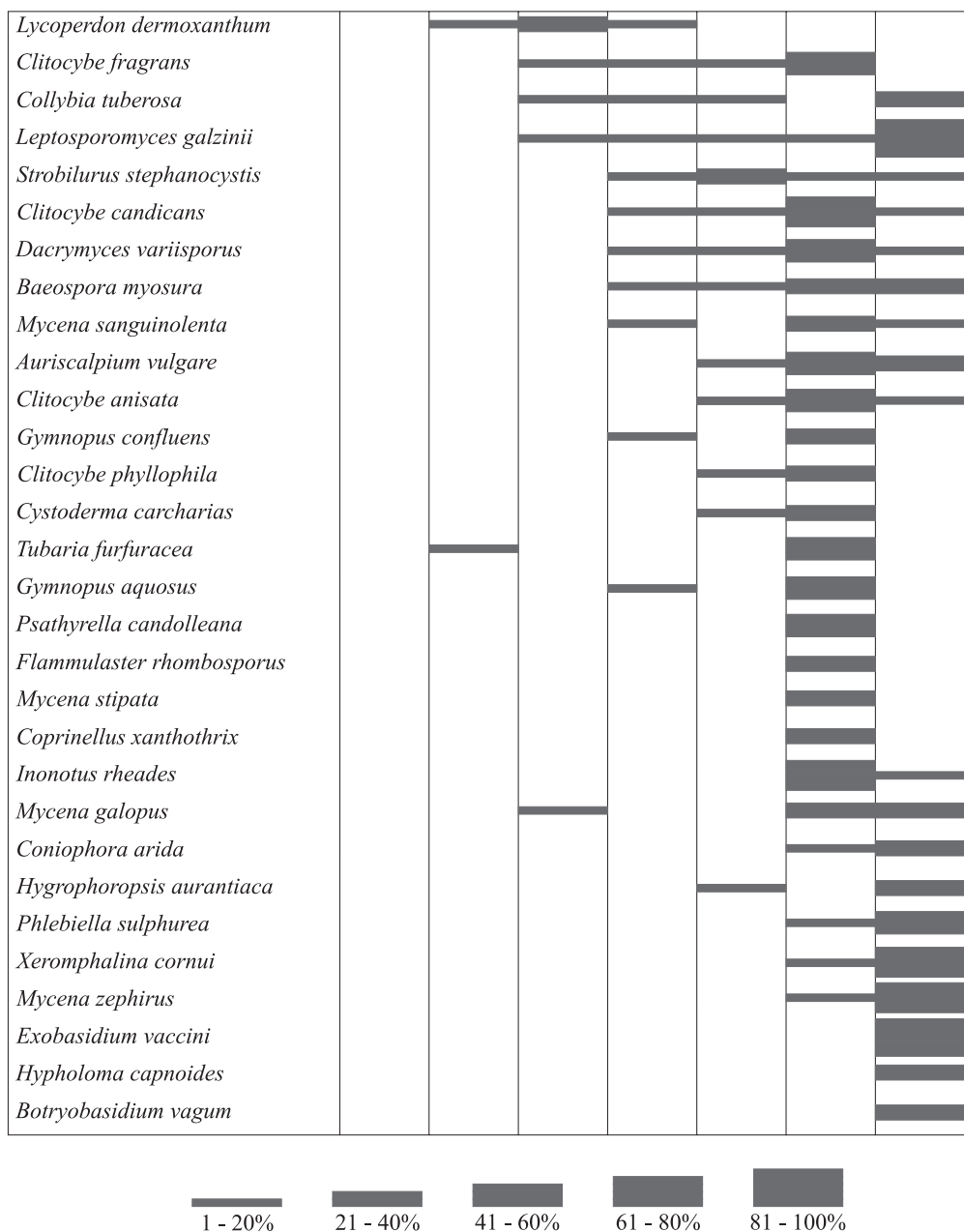


Fig. 26. Spatial frequency of most frequent non-ECM fungi in consecutive phases of the spontaneous secondary succession on abandoned farmland.

instance *Amanita muscaria*, *Chalciporus piperatus*, *Tricholoma equestre* and *Boletus edulis*, after attaining maximum abundance a decrease in carpophore production is not immediately followed by decreasing spatial frequency – the frequency tends to stay high longer. Similarly, high carpophore abundances combined with high spatial frequencies typical of *Inocybe lacera*, *Suillus luteus*, *Amanita muscaria* and *Laccaria proxima* early after trees encroachment suggest rapid and simultaneous establishment of numerous genets of these species originating from spore dispersal.

#### Changes in species composition of ECM fungi

An open community of psammophilous grassland with 6-8(12) year old pines was characterized by the presence of a rich and diverse community of ECM fungi corresponding to the ‘initial floristics’ model (EGLER 1954). Among them a small group of species attained not only very high spatial frequency but also high sporocarp abundance, which suggests that they reveal a special affinity for the youngest trees and might have started fruiting before the other species. In this group are *Inocybe lacera*, *Laccaria proxima*, *Suillus luteus*, *Cortinarius cinnamomeoluteus* and *Amanita muscaria*. These species also accounted for most of the sporocarp biomass production in early succession phases. At least the first three species are regarded as ‘early stage fungi’ (MASON *et al.* 1982, 1983; DIGHTON & MASON 1985). However, they were not soon replaced by other species but persisted in developing vegetation at least till the pre-forest phase – the juniper-aspen brushwood.

The majority of the ECM species present in the initial, optimal and early terminal stages of the old-field succession are rather ‘multi-stage fungi’, similar to the observations in jack pine stands regenerating after a wildfire (VISSER 1995) or in primary succession in a glacier forefront (JUMPPONEN *et al.* 2002). However, most of these ‘multi-stage fungi’ showed preferences for a particular period of vegetation development, which was manifested with more abundant sporocarp production and/or higher spatial frequency.

In the terminal pine forest, species regarded as ‘late stage fungi’, especially of the genus *Russula*, but also *Amanita*, *Cortinarius*, *Lactarius* and *Xerocomus* prevailed. Most of the typical forest species were recorded in the successional series late, only in the nearly mature stand. Relatively few ‘late stage species’ started fruiting earlier: *Lactarius rufus* and *Xerocomus badius* in the stands around canopy closure, and *Amanita citrina*, *Cantharellus cibarius* and *Russula paludosa* in the juniper-aspen brushwood. It seems that, in contrast to the initial rapid and somewhat chaotic species colonization, the recruitment of new species along with advancing age and maturity of the forest long after the canopy closure is much slower and gradual.

The pattern described above seems to agree well with the ‘early-’ and ‘late-stage’ model, especially if it is amended by adding ‘multi-stage fungi’. The model has played a very inspiring role in mycorrhizal ecology and research on spatio-temporal changes in ECM communities. It was supported (e.g., JANSEN 1991; VISSER 1995; HELM *et al.* 1996) as well as opposed (e.g., TERMORSHUIZEN 1991; KEIZER & ARNOLDS 1994; JUMPPONEN *et al.* 1999; BONET *et al.* 2004). In the present study, an occurrence of a narrow group of species specific for the intermediate part of the chronosequence (pines 10-20 years old, the cover exceeding 50%), namely *Cortinarius anthracinus*, *C. colus*, *C. mucosus*, *C. paraevernius* and numerous species of *Tricholoma* cannot be explained within the early-, late- or multi-stage fungi concepts unless they were regarded as intermediate-stage fungi. Some of these species were present also in the middle-aged pine stands described, e.g., by HINTIKKA (1988; oligotrophic, mainly 20-30 and 30-50 years old stands, 45-90% canopy closure) and TERMORSHUIZEN

(1991; spontaneous, 8-13 years old stand on drifting sands, 85-95% canopy closure). An abundant occurrence of carpophores of different species of *Tricholoma* in pine stands around canopy closure was observed also on mineral soils with no humus yet formed, in pine plantations on post-arable land and on post-mining heap (KAŁUCKA, unpubl. data).

Changes of the ECM fungi composition in the chronosequence studied can be at least partly explained as the sequence of fungi representing different life histories. Various assemblages of species found in stands of different ages are believed to exhibit characters that are adaptations to changes in habitat conditions as the trees mature (FRANKLAND 1998). In the earlier succession phases, r-selected species prevailed, growing on mineral soil, producing relatively small carpophores but in mass quantities, such as *Inocybe lacera*, *Cortinarius cinnamomeoluteus* and other species of this genus with tiny fruit bodies, *Hygrophorus hypothejus* and *Laccaria proxima*. Later, on xeric mineral soil with low C and N content and high row litter cover, species producing abundant, but bigger carpophores became numerous, mainly *Tricholoma* spp. and some *Cortinarius* spp. (stress-tolerant species?). In the pine forest, with developing organic layers, big carpophores of different *Russula* species as well as *Cortinarius* (mainly *C. caperatus*) and *Amanita* prevailed, but their abundance is much lower; such species are usually regarded as K-selected (DIGHTON & MASON 1985; LAST *et al.* 1987).

Strong links between the patterns of the ECM species composition, frequency and abundance, and vegetation development, age of trees, stand structure and soil development suggest that ECM fungal succession would be better explained against a background of vegetation succession and stages of stand development. It seems that the successional trajectory of ECM fungi of the spontaneous, old-field series leading to the formation of fresh pine forest is a 3-stage process:

- colonization stage – rapid colonization by a rich ECM community dominated by *Inocybe lacera*, *Laccaria proxima*, *Suillus luteus*, *Cortinarius cinnamomeoluteus*, *Amanita muscaria* and *Hygrophorus hypothejus*; increase in species richness and diversity, sporocarp abundance and biomass production (phases 4 and 5, initial and early optimal stage of vegetation development: open psammophilous grassland with junipers and pines up to 12(16) years old);
- stabilisation stage – stabilisation of the ECM community, prevalence of *Laccaria proxima*, *Suillus luteus*, *Cortinarius cinnamomeoluteus* and other small-sized *Cortinarius* species, *Amanita muscaria*, *Hygrophorus hypothejus* and *Chalciporus piperatus*, arriving and abundant occurrence of *Lactarius rufus*, *Cortinarius mucosus* and *C. parevernius*, more and more abundant occurrence of many species of *Tricholoma*; further increase in the species richness and diversity, maximum overall carpophore abundance and biomass production (phases 6 and 6/7; advanced and late optimal stage of vegetation development: psammophilous grassland with rich moss and lichen layer, juniper shrubs and pines up to 18(25) years old, ca 50% pine cover up to canopy closure); the period of canopy closure with intensified litter accumulation and xeric soil conditions marked with decreasing carpophore numbers and prevalence of *Tricholoma* spp. in the biomass production;
- replacement stage – gradual change in the ECM species composition, partial replacement of species typical of the pre-forest vegetation and arrival of numerous forest species, mainly of *Russula*, *Lactarius*, *Cortinarius* and *Amanita*; increase in the species diversity, decrease in the species richness, conspicuous decrease in the overall carpophore abundance and biomass, prevalence of *Russula* in the biomass production (phases 8 to

10, terminal stage of vegetation development; formation of the forest vertical structure, development of soil profile and humus layer, pines over 30 years old, up to 100 in the present study); if the vegetation successional trajectory comprises a transitional, early terminal community of the juniper-aspen brushwood, then a presence of numerous species host-specific to deciduous trees is considerable resulting in a temporal shift in species richness and prevalence of *Cortinarius* in the carpophore biomass production.

A similar, 3-stage scheme of the ECM fungi succession (based on the occurrence of ECM roots) was proposed by TWIEG *et al.* (2007) for the successional series of mixed, Douglas fir and paper birch temperate forest developing after stand-replacing fire and clearcutting; a kind of classification of ECM fungi based on their presence in different stages of forest development was also proposed by KEIZER & ARNOLDS (1994). It seems that some common general patterns could influence the ECM fungal succession during secondary succession of the forest community.

### Dynamics of ectomycorrhizal roots

As summarized by SMITH & READ (2008), fine roots of trees (< 2 mm in diameter) are predominantly found in the superficial organic layers and most of them in boreal and temperate forest biomes are ectomycorrhizal. The fine roots of Norway spruce in European forests develop mainly in the upper 10 cm of soil (STOBER *et al.* 2000). Most fine root biomass (over 80%) of Scots pine was found in the upper (0-15 cm) soil level (OLEKSYN *et al.* 1999). A similar distribution of *Pinus sylvestris* ECM root biomass was also found by PERSSON (1983). As much as 80% of ectomycorrhizal tips can occur in the top 5 cm of soil profile (MIKOLA *et al.* 1966). In boreal *Pinus sylvestris*-*Picea abies* forests, ECM roots most intensively penetrate the superficial organic layers of the podzol, but numerous mycorrhizae of various species occur also in the mineral horizons (ROSLING *et al.* 2003).

Concentration of the fine roots, of which more than 90% were always ectomycorrhizal, in the superficial soil part was observed also in the chronosequence studied. However, the main body of the roots moved from the mineral to organic horizons along with the vegetation succession and development of the soil profile. A similar trend was found by VOGT *et al.* (1981) in stands of *Abies amabilis* in the USA. Accumulating litter and humus layer formed underneath are the substrates, which can be intensively exploited at least by some of the ECM species associated with mature stands as a source of nitrogen (SMITH & READ 2008).

The length of the fine roots and the number of ECM tips found in the subsequent succession phases did not differ significantly apart from phase 8 (juniper-aspen brushwood), where they were much higher. However, the average length of the roots as well as the number of ectomycorrhizae in the initial and optimal stage of vegetation development (pines up to 20-25 years old) were usually higher than in the pine forest (pines ca 100 years old). Similar observations were made by TERMORSHUIZEN (1991) in *Pinus sylvestris* stands of different ages in the Netherlands. In the younger stands, 4-13 years old, the length of ECM roots and the number of ECM tips were usually higher than in the older stands, 50-80 years old, and standard deviations were very high. JANSEN (1991) observed a distinct decrease in the number of ECM tips from the average 128/100 cm<sup>3</sup> in young stands (8-18 years old) to 30/100 cm<sup>3</sup> in older stands (41 and 54 years old) in the plantations of *Pseudotsuga menziesii* in the Netherlands. In the *Pseudotsuga menziesii* stands, the culmination of the fine root biomass and ectomycorrhizae coincides with attaining maximum biomass of foliage and crowns (VOGT *et al.* (1983). Later the biomass decreases, earlier and faster in the rich sites than in the poor sites.



An occurrence of numerous fine roots and ectomycorrhizae may be favoured by rather thin humus and sparse field layer. TERMORSHUIZEN (1991) found that in 60 years old pine stands characterized by such features the number of ECM root tips and the length of fine roots were as high as in young stands.

An exceptionally high density of tree roots in the juniper-aspen brushwood may be an effect of mixed character of the stand and the predominance of deciduous trees with high proportion of aspen (STRONG & LA ROI 1983; VILLENEUVE *et al.* 1989; TWIEG *et al.* 2007). STRONG & LA ROI (1983) compared the stands of *Pinus banksiana*, *Populus tremuloides* and *Picea glauca* in different age and on different soils and they found that in the 48 year old aspen stands on mineral sandy soil the densities of roots, both total and in the superficial soil, were the highest. The roots were concentrated mainly in the zone between the organic and mineral layers.

In all the succession phases studied, independently of the stand composition, age of trees and high standard deviations in the root length and the number of mycorrhizas, high positive correlation was found between the length of living fine roots and the number of ECM tips. Also, the ramification index, i.e., the number of ECM tips per cm of the root length, shows no statistically significant differences among phases in a given year and is characterized with much smaller standard deviations. In two seasons it attained the values of 4.7-5.7 and 7.0-8.9, respectively. In *Pinus sylvestris* stands in different ages and of different origin (spontaneous and planted, different rotations, different soils) in the Netherlands, TERMORSHUIZEN (1991) found a similar correlation, but the ramification index was 1.6-2.0. According to MEYER (1987), the index is an indication of the fine root vitality. In the *Pinus sylvestris* stands situated in a zone of low air pollution in Poland, the ramification index equaled 2.7-3.8, and in the stands suffering from industrial emissions it was 1.3-2.2 (JÓZEFACIUKOWA *et al.* 1995). High indices recorded in the old-field chronosequence studied may indicate good condition of trees and high mycorrhizal activity of their roots.

## 7. SUMMARY OF RESULTS AND CONCLUSIONS

1. The research on the macrofungi in the process of spontaneous secondary succession was carried out in the chronosequence of old fields abandoned at different times and inhabited by vegetation at different succession stages. The study area is situated near the southwest peripheries of the Białowieża old-growth forest (NE Poland), in two nature reserves, Jelonka and Czechy Orłańskie. Both sites are extensively studied in the framework of long-term ecological research on the process of natural fresh coniferous forest return after centuries of deforestation and agricultural management conducted by the Białowieża Geobotanical Station, University of Warsaw.
2. The sporocarps of macromycetes were subject to qualitative and quantitative analyses in three consecutive seasons (17 field observations); the study focused on basidiomycetes as the proportion of ascomycete fruit bodies was negligible. The ECM roots were also examined in two seasons for the number of ECM tips and fine root length. A series of soil analyses, as well as vegetation examination, were performed to document concurrent habitat features.

3. The succession of macrofungi was examined against the background of the old-field succession model proposed by FALIŃSKI (1986b) and FALIŃSKI *et al.* (1993). Eleven permanent plots, 1000 m<sup>2</sup> each, situated in the Jelonka reserve, represented 6 phases of secondary succession, from a fallow herb-grassland (phase 2), a complex of psammophilous grasslands, juniper shrubs and Scots pine thickets (phases 4-6/7) to a brush-wood of juniper, trembling aspen and Scots pine (phase 8), corresponding to the initial, optimal and early terminal stage of vegetation development. Five plots representing nearly mature pine forest of *Peucedano-Pinetum* type (phase 9/10, pines ca 100 years old), situated in the Czechy Orlańskie reserve were also included, each of them of 400 m<sup>2</sup>. Each succession phase studied was represented by a total area of 2000 m<sup>2</sup>, with the exception of the earliest phase 2, for which one 1000 m<sup>2</sup> plot was selected.
4. The series yielded 300 species of macromycetes (including 6 collective taxa) found in permanent plots. Among them were 98 species of ECM fungi (32.7%), 200 species of saprotrophic fungi (66.7%) and 2 species of plant parasites (0.7%). Additional 18 species were found outside the plots.
5. In the early successional, tree-less herb-grassland (phase 2, initial stage) 15 basidiomycete species were found. All the species recorded were saprotrophic, growing on dead plant remnants or on soil. The highest spatial frequency and abundance was achieved by *Crinipellis scabellia*, *Marasmius oreades*, *Mycena olivaceomarginata*, *M. flavoalba* and *M. leptcephala*, *Coprinopsis friesii* and *Galerina* spp.
6. Along with the development of poor psammophilous grassland and colonization of the area by the ECM trees, rapid establishment of macrofungi occurred; in phase 4 (initial stage), with most of pines 6-8 (up to 12) years old, a total of 92 species were recorded. Among them 44 species were ectomycorrhizal, most of them host-specific to *Pinus sylvestris*. The group of species comprising *Inocybe lacera*, *Suillus luteus*, *Amanita muscaria*, *Cortinarius cinnamomeoluteus*, *Laccaria proxima* and *Hygrophorus hypothejus* prevailed in terms of spatial frequency and carpophore abundance. The highest proportion of the carpophore biomass production was achieved by the genera *Amanita*, *Suillus* and *Inocybe*. Among saprotrophic fungi, species growing on soil and among mosses prevailed.
7. In the optimal stage of vegetation development, changes in the fungal community, especially in the ECM fungi, were mainly of quantitative character. In open complex of psammophilous grassland with rich lichen-moss bottom layer, juniper shrubs and 8-15 (up to 20) years old pine thickets (phase 6), a total of 114 macromycete species were found. Among them were 47 ECM species; with an exception of *Inocybe lacera*, the group of species listed earlier still prevailed, joined by *Chalciporus piperatus*, *Laccaria bicolor*, *Lactarius rufus* and a few species of *Tricholoma*. The presence of *Inocybe curvipes*, *Cortinarius parevernus*, *C. anthracinus*, *C. mucosus*, *C. colus*, some other *Cortinarius* species with rather small carpophores and *Sacrodon squamosus* was also specific for this stage of vegetation development. The end of the optimal stage (pines 12-18 up to 25 years old), characterised by progressing canopy closure and xeric soil conditions, was marked by the presence of numerous species of *Tricholoma* and their exceptionally high proportion in the carpophore biomass production. A part of the ECM species present in the optimal stage of the vegetation succession did not continue fruiting after the canopy closure. Only two new species which continued fruiting in the pine forest arrived, *Lactarius rufus* and *Xerocomus badius*. Saprotrophic species growing on soil and among mosses were still numerous, but the diversity and abundance of litter- and wood-inhabiting species conspicuously increased.

8. The fungal species richness in the juniper-aspen brushwood (phase 8, early terminal stage) increased to 183 species, 70% of which were saprotrophic ones. Litter- and wood-inhabiting fungi prevailed, while the species typical of open vegetation decreased in number, frequency and abundance. The ECM species composition was relatively similar to earlier phases, but the carpophore abundance and productivity decreased. Very high proportion of aspen, birch and willow in the stand resulted in the increased number of ECM species host-specific to deciduous trees, making the ECM species richness very high (49 species). In terms of carpophore biomass, *Amanita* and *Cortinarius* prevailed. Relatively few ECM species typical of forest communities arrived at this stage of vegetation development. These were *Amanita citrina*, *Cantharellus cibarius* and *Russula paludosa*.
9. In the fresh pine forest *Peucedano-Pinetum*, being the final community of the successional series studied, the fungal species richness decreased to 30 ECM and 68 non-ECM fungi. The abundance and overall carpophore biomass was also the lowest. However, in comparison with the optimal stage of vegetation development, the species diversity, both in the ECM and saprotrophic species was higher. The species composition of ECM fungi changed considerably; many species present in earlier vegetation phases were replaced by species typical of forests, such as species of *Russula*, *Cortinarius*, *Lactarius* and *Amanita*. *Russula* was the most productive genus, together with *Cortinarius* they accounted for about 60% of the carpophore biomass production. Among the saprotrophic fungi, species growing on litter and wood prevailed; the majority of exclusive saprotrophic species occurring in the forest were wood-inhabiting taxa.
10. The species colonization pattern corresponds to the 'initial floristics' model, especially for the ECM fungi. A rapid initial increase in the species number was observed, followed by long persistence and changing prevalence of different groups of species over time. The majority of the ECM species can be regarded as early arriving multistage fungi. Among them a group of species occurring along the whole chronosequence where trees were present (phases 4-9/10) can be distinguished as well as groups of species which stop fruiting after canopy closure or before the mature forest formation. The species richness is built primarily by an accumulation of species, with a distinct although not numerous group of species specific for the optimal stage of vegetation development, followed by partial but considerable replacement best seen between the brushwood and forest phases of the terminal succession stage. When the forest community mature, gradual recruitment of new species can be observed.
11. The majority of the ECM species present in the initial, optimal and early terminal stage of the old-field succession can be regarded as 'multi-stage fungi'. However, in early succession phases species regarded as 'early stage fungi' prevailed in terms of spatial distribution, carpophore abundance and biomass production, such as *Inocybe lacera*, *Laccaria proxima*, *Suillus luteus*, and also *Cortinarius cinnamomeoluteus* and *Amanita muscaria*. They were not soon replaced by other species but persisted in developing vegetation at least till the pre-forest phase – the juniper-aspen brushwood. In the pine forest, species regarded as 'late stage fungi', especially of the genus *Russula*, but also *Amanita*, *Cortinarius*, *Lactarius* and *Xerocomus* prevailed. This pattern can also be explained in terms of changing proportions of fungi representing different life histories – from r- to K-selected species.
12. A three-stage process for the succession of ECM fungi in the poor post-arable land till the formation of mature pine forest community is proposed:

- colonization stage – rapid colonization by a rich ECM community dominated by *Inocybe lacera*, *Laccaria proxima*, *Suillus luteus*, *Cortinarius cinnamomeoluteus*, *Amanita muscaria* and *Hygrophorus hypothejus*; fast increase in the species richness and diversity, sporocarp abundance and biomass production;
  - stabilisation stage – stabilisation of the ECM community, prevalence of *Laccaria proxima*, *Suillus luteus*, *Cortinarius cinnamomeoluteus* and other small-sized *Cortinarius* species, *Amanita muscaria*, *Hygrophorus hypothejus* and *Chalciporus piperatus* joined with abundant occurrence of *Lactarius rufus*, *Cortinarius mucosus*, *C. parevernus* and many species of *Tricholoma*; further but slower increase in the species richness and diversity, maximum overall carpophore abundance and biomass production. Late in this stage *Tricholoma* spp. prevail in the biomass production, abundance of carpophores of other species decreases. This stage coincides with the period of reaching stand canopy closure;
  - replacement stage – gradual change in the ECM species composition, partial replacement of species typical of the pre-forest vegetation and establishment of numerous forest species, mainly of *Russula*, *Lactarius*, *Cortinarius* and *Amanita*; increase in the species diversity, decrease in the species richness, conspicuous decrease in the overall carpophore abundance and biomass, prevalence of *Russula* in the biomass production.
13. Changes in the fungal species composition as well as patterns of the carpophore spatial frequency, abundance and biomass production are strongly linked to the vegetation development, its rate, age of trees, stand composition and structure and soil development. Succession of the ECM fungi in the studied coniferous series seems to be affected mainly by the age of trees, stand structure and development of soil profile, but the species composition (and usually not fruiting dynamics) can be modified by the presence of additional, deciduous tree species and the presence of older and successional more advanced vegetation stages in close vicinity. Succession of the saprotrophic species seems to be primarily affected by the presence and diversification of available substrates and microclimatic conditions resulting from the plant community composition, stand composition and structure and accumulation of organic debris.
  14. The ECM roots were located in the mineral superficial soil in earlier succession phases. Later, along with the soil development, they tended to concentrate in between the mineral soil and accumulating litter and finally in the organic horizons in the terminal stage of vegetation succession. The length of fine roots and the number of ECM tips were the highest in the juniper-aspen brushwood as a result of a high proportion of aspen. Independently of the stand composition, age of trees and high standard deviations in the root length and the number of mycorrhizae, high positive correlation was found between the length of living fine roots and the number of ECM tips. The ramification index showed no statistically significant differences among phases in a given year and was characterized with much smaller standard deviations. Its relatively high values may indicate good condition of trees and high mycorrhizal activity of their roots.
  15. The present study confirmed strong links and relationships between the succession of fungi and vegetation; the two processes parallel each other, and the species composition and dynamics of carpophore production of fungal assemblages match well with successional changes in phytocoenoses. Without the analysis of vegetation successional trajectory, interpretation of fungal succession, including both ECM and saprotrophic species, would be incomplete.

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**Dr. Izabela Kałucka**  
 Department of Mycology  
 University of Łódź  
**Banacha 12/16**  
**PL-90-237 Łódź**  
 ikałucka@biol.uni.lodz.pl

## Appendix 1

### Soil analysis in the permanent plots

#### Descriptions of the soil profiles

Phases 2-8 – Jelonka reserve, post-arable rusty soil. Phase 9/10 – Czechy Orlańskie reserve, podzolic soil developing on post-arable rusty soil. Soil profiles down to ca 60 cm.

##### Initial stage:

Phase 2 – plot 82		
Ap	0-31 cm	Plough humous horizon with abrupt boundary; loamy sand, dark brown, with stones of a few centimetres in diam. at places. Earthworms present.
Bv	> 31 cm	Sand of rusty brown colour with considerable amount of gravel and stones.
Phase 4 – plot 84		
Ap	0-28 cm	Plough humous horizon with abrupt boundary; loamy sand of dark brown grey colour, solitary stones at places. Concentration of root biomass.
Bv	28-68 cm	Loamy sand of light brown colour with considerable amount of gravel and stones of a few centimetres in diam.

##### Optimal stage:

Phase 5 – plot 61		
Ap	0-24 cm	Plough humous horizon with abrupt wavy boundary; loamy sand of dark grey colour. Concentration of root biomass.
Bv	> 24 cm	Loamy sand of pale yellowish colour. Considerable amount of stones distributed at the depth of ca 40 cm.
Phase 5/6 – plot 13		
Ap	0-25 cm	Plough humous horizon with abrupt boundary; loamy sand of dark grey colour. Concentration of root biomass.
Bv	> 25 cm	Loamy sand of pale yellowish colour. Considerable amount of stones distributed at the depth of ca 50 cm.
Phase 6 – plot 1		
Ap	0-30 cm	Plough humous horizon; loamy sand of brown grey colour, intensely overgrown with plant roots. The layer of stones of a few to several centimetres in diam. present in the contact zone with horizon Bv.
Bv	30-54 cm	Loamy sand, rusty yellow. Layers of loamy sand of rusty colour at places. Solitary stones of a few centimeters in diam. in the whole horizon. Roots evenly distributed.
Phase 6 – plot 6		
Ap	0-20 cm	Plough humous horizon with abrupt boundary; weakly loamy sand of dark grey colour, overgrown with roots.
Bv	> 20 cm	Sand of rusty grey yellow colour, with relatively few tree roots at places. Humic wedges after roots decay.

**Terminal stage:**

## Phase 8 – plot 16

Ol	1-0 cm	Litter horizon consisting of leaves of aspen and birch, juniper needles, solitary pine needles and dead grass, loose. Differentiation into the horizons of Ol and Of can be seen at places.
A	0-1.5 cm	Humus horizon of initial mull type or initial moder in hollows; weakly loamy sand of black grey colour, intensely overgrown with mycelium and roots. In the places where pine or spruce grow, solitary bleached quartz grains are present (initiation of the podzolization process).
Ap	1.5-25 cm	Plough humous horizon with abrupt boundary; loamy sand of grey brown colour, overgrown with roots of herbaceous plants and trees.
Bv	25-53 cm	Loamy sand of yellow rusty colour, rusty at places, with solitary roots of trees. Irregular lower boundary.
Bv/C	> 58 cm	Sand of yellow colour with clay layers of rusty colour.

## Phase 9/10 – plot 6C

Ol	6-5 cm	Litter horizon consisting of pine needles, pieces of bark and fallen twigs.
Of	5-2 cm	Fermentation horizon; the material partly fragmented, the structure partly preserved. Mainly pine needles and twigs, overgrown with mycelium and roots.
Oh	2-0 cm	Epilumus with solitary bleached quartz grains. Humus type – mor.
AEes	0-1.5 cm	Eluvial horizon; sand of light pale colour, with bleached quartz grains prevailing. Sparse roots present.
Bfe	1.5-20 cm	Illuvial horizon (with iron); sand of rusty colour, gradually changing into Ap.
Ap	20-35 cm	Plough humous horizon with abrupt boundary; loamy sand of grey pale colour, with roots and solitary humic wedges after roots decay.
Bv	> 35 cm	Fine, loamy sand of yellow pale colour.

Table A1.1  
Texture of the soil along the studied successional gradient

Succession stage	Succession phase	Plot no.	Soil horizon	Skeleton parts [mm]				Earth parts [mm]								Total sand	Total silt	Total clay	Earth parts in sample	Textural class	
				total	stones >20	coarse gravel 20-5	fine gravel 5-2 2-1	sand		silt		clay									
								1-0.5	0.5-0.25	0.10-0.05	0.05-0.02	0.02-0.006	0.006-0.002	<0.002							
															% in earth parts						
				% in sample				% in earth parts								% %					
Initial	2	82	Ap	29.68		6.16	11.0	12.52	17.5	32.25	23.25	9	8	3	4	3	73	17	10	70.32	w.l.s.
			Bv	62.47	25.0	9.68	11.28	16.51	33.15	24.50	14.35	7	7	5	3	6	72	14	14	37.53	l.l.s.
	4	84	Ap	15.50		1.67	4.59	9.24	11.35	37.50	24.15	9	6	5	3	4	73	15	12	84.50	l.l.s.
			Bv	15.0		2.49	5.28	7.23	18.0	40.9	24.1	5	2	4	2	4	83	7	10	85.0	w.l.s.
Optimal	5	61	Ap	7.29			3.32	3.97	10.0	41.25	27.75	8	4	4	2	3	79	12	9	92.71	w.l.s..
			Bv	5.09			2.0	3.09	9.3	37.75	34.95	7	3	3	2	3	82	10	8	94.91	w.l.s.
	6	13	Ap	5.55			2.99	2.56	8.6	39.65	30.75	7	4	4	3	3	79	11	10	94.45	w.l.s.
			Bv	2.99			1.38	1.61	4.45	39.45	35.1	8	5	4	2	2	79	13	8	97.01	w.l.s.
Terminal	6/7	1	Ap	4.14			1.62	2.52	10.85	41.50	31.65	7	3	2	1	3	84	10	6	95.86	w.l.s.
			Bv	36.02	16.38	10.49	5.66	3.49	10.7	31.9	30.4	11	7	4	2	3	73	18	9	63.98	w.l.s.
	6/7	6	Ap	3.74			1.33	2.41	7.50	42.05	36.45	5	3	2	1	3	86	8	6	96.26	w.l.s.
			Bv	1.66				1.66	3.95	41.65	40.40	7	3	2	1	1	86	10	4	98.34	l.s.
Terminal	8	16	Ap	9.01		1.60	2.56	4.85	8.50	43.95	26.55	7	5	4	1	4	79	12	9	90.99	w.l.s.
			Bv	4.24			1.58	2.66	5.0	40.75	30.25	13	3	4	2	2	76	16	8	95.76	w.l.s.
	9/10	6C	AEes	4.58			1.37	3.21	10.50	39.30	35.20	6	4	2	0	3	85	10	5	95.42	l.s.
			Bfe	4.18			1.44	2.74	8.45	41.9	36.65	5	3	2	0	3	87	8	5	95.82	l.s.
Terminal			Ap	3.69			1.21	2.48	9.8	40.75	32.45	8	1	2	1	5	83	9	8	96.31	w.l.s.
			Bv	3.50				3.50	8.1	38.35	34.55	8	5	1	1	4	81	13	6	96.50	w.l.s.

l.s. – loose sand; w.l.s. – weakly loamy sand; l.l.s. – light loamy sand

Table A1.2  
Some chemical properties of the soil along the studied successional gradient – soil profiles

Succession stage	Succession phase	Plot no.	Soil horizon	pH		C %	N %	C/N
				in H <sub>2</sub> O	in KCl			
Initial	2	82	Ap	5.5	4.4	1.39	0.093	14.9
			Bv	5.7	4.6	0.381	0.033	11.5
	4	84	Ol	4.9	4.4	32.2	1.27	25.4
			Ap	5.1	4.3	1.16	0.096	12.1
			Bv	5.6	4.4	0.068	0.011	6.2
Optimal	5	61	Ol	4.5	3.9	31.8	0.695	45.8
			Ap	4.9	4.3	0.831	0.056	15.2
			Bv	4.9	4.5	0.183	0.019	9.6
	6	13	Ol	4.8	4.2	30.3	1.18	25.7
			Ap	4.8	4.3	0.997	0.062	16.1
			Bv	4.9	4.5	0.305	0.021	14.5
	6/7	6	Ol	4.9	4.4	32.2	1.04	31.0
			Ap	4.9	4.4	0.550	0.031	17.7
			Bv	4.8	4.6	0.232	0.020	11.6
Terminal	8	16	Ol	6.1	5.8	34.4	1.70	20.2
			Of	5.9	5.5	22.8	1.55	14.7
			A	5.2	4.5	5.67	0.442	12.8
			Ap	5.2	4.4	0.595	0.033	18.0
			Bv	4.9	4.6	0.194	0.016	12.1
	9/10	6C	Ol	4.3	3.8	35.2	0.607	58.0
			Of	4.0	3.2	31.3	1.57	19.8
			Oh	3.9	3.1	20.3	1.10	18.4
			AEes	4.1	3.2	0.922	0.045	20.5
			Bfe	4.8	4.2	0.700	0.037	18.9
			Ap	5.0	4.4	0.543	0.029	18.7
			Bv	5.0	4.5	0.260	0.017	15.3



Table A1.3

Some chemical properties of the soil along the studied successional gradient – bulk surface soil (0-15 cm)

Succession stage	Succession phase	Plot no.	pH		C %	N %	C/N
			in H <sub>2</sub> O	in KCl			
Initial	2	82	5.5	4.8	1.63	0.113	14.4
	4	67	4.6	4.1	1.43	0.089	16.1
		84	4.8	4.2	1.16	0.080	14.5
Optimal	5	61	4.8	4.2	1.25	0.077	16.2
		85	4.8	4.2	1.08	0.073	14.8
	6	7	4.7	4.2	1.05	0.061	17.2
		13	4.8	4.2	1.07	0.058	18.4
	6/7	1	4.8	4.2	0.957	0.054	17.7
		6	4.8	4.2	0.833	0.053	15.7
Terminal	8	11	5.3	4.5	1.20	0.057	21.0
		16	5.0	4.2	1.07	0.060	17.8
	9/10	2C-8C	4.6	4.1	0.983	0.041	24.0

Table A1.4

Surface soil moisture (%) in observation plots compared among sampling terms  
(26/07, 09/09, 12/09, 20/10, 06/11)

Succession stage	Succession phase	Plot no.	Minimum median (sampling term)	Maximum median (sampling term)	Kruskal-Wallis test df=4
Initial	2	82	9.67 (09/09/)	18.15 (20/10)	n=30; H=10.00*
	4	67	6.88 (09/09/)	14.04 (06/11)	n=30; H=14.90**
		84	5.02 (09/09/)	14.85 (06/11)	n=29; H=15.51**
Optimal	5	61	4.76 (09/09/)	14.91 (06/11)	n=30; H=21.13***
		85	3.78 (09/09/)	12.37 (06/11)	n=30; H=12.38*
	6	7	5.73 (09/09/)	13.53 (20/10)	n=29; H=15.31**
		13	6.00 (09/09/)	10.64 (06/11)	n=30; H=17.89**
	6/7	1	4.24 (09/09/)	10.20 (06/11)	n=30; H=20.08***
		6	4.00 (09/09/)	15.18 (06/11)	n=30; H=12.67*
Terminal	8	11	7.12 (09/09/)	17.89 (06/11)	n=30; H=11.88*
		16	6.66 (09/09/)	20.10 (06/11)	n=30; H=18.35**
	9/10	2C, 5C, 6C, 7C, 8C	11.05 (09/09/)	24.54 (26/07)	n=60; H=6.69 <sup>ns</sup>

ns – not significant differences,  $p > 0.05$ ; significant difference for at least one pair of sampling terms at \*  $p < 0.05$ , \*\*  $p < 0.01$ , \*\*\*  $p < 0.001$

Table A1.5  
Surface soil moisture (%) in observation plots compared among the plots  
(82 – phase 2; 67, 84 – phase 4; 61, 85 – phase 5; 7, 13 – phase 6; 1, 6 – phase 6/7;  
2C, 5C, 6C, 7C, 8C – phase 9/10)

Sampling term	Minimum median (plot no.)	Maximum median (plot no.)	Kruskal-Wallis test df=11
26/07	7.82 (1)	24.54 (2C, 5C, 6C, 7C, 8C)	n=128; H=74.64****
09/09	3.78 (85)	11.05 (2C, 5C, 6C, 7C, 8C)	n=65; H=23.94*
12/09	4.79 (13)	14.59 (2C, 5C, 6C, 7C, 8C)	n=65; H=44.78****
20/10	7.46 (6)	20.54 (2C, 5C, 6C, 7C, 8C)	n=65; H=31.20**
06/11	10.20 (1)	23.96 (2C, 5C, 6C, 7C, 8C)	n=65; H=34.01***

significant difference for at least one pair of plots at \*  $p < 0.05$ , \*\*  $p < 0.01$ , \*\*\*  $p < 0.001$ , \*\*\*\*  $p < 0.0001$

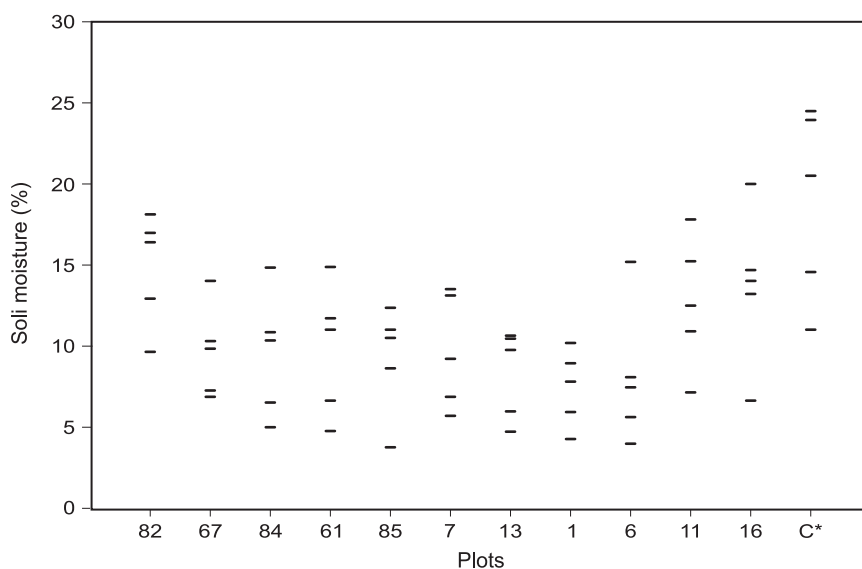


Fig. A1.1. Surface soil moisture (%) in permanent plots – medians in five sampling terms (26/07, 09/09, 12/09, 20/10, 06/11); for numbers of samples see 4. Material and methods.

C\* – stands for the plots 2C, 5C, 6C, 7C, 8C (phase 9/10); soil samples from these plots have been analysed together.

## **Appendix 2**

### **Vegetation description in the permanent plots**

Table A2.1

Vegetation description in the permanent plots in the Jelonka reserve

(June 1997; Londo method; [ ] – cover by the crowns of trees growing outside the examined subplot; n – no data)

Succession stage		Initial																								
Succession phase		2					4										5									
Number of plot		82					67					84					61									
Subplot		a	b	c	d	e	a	b	c	d	e	a	b	c	d	e	a	b	c	d	e	a	b			
Subplot area (m²)		25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	25	
Cover of plant layers (%)	a	·	·	·	·	·	·	·	·	·	·	10	60	·	·	·	[10]	·	30	10	·	5	[5]			
	b	<5	·	·	·	·	·	60	10	30	25	20	20	10	20	10	10	30	10	30	10	40	50			
	c	100	100	100	100	100	95	80	90	30	50	60	40	10	50	20	40	50	80	20	90	70	40			
	d	20	10	10	10	10	50	60	85	100	100	70	50	90	80	60	80	80	40	90	50	45	60			
	d <sub>B</sub>	20	10	10	10	10	30	50	60	90	90	60	50	80	80	40	40	30	30	50	20	30	40			
including mosses	d <sub>L</sub>	·	·	·	·	·	20	10	25	30	10	10	5	10	<5	20	40	50	10	40	30	15	20			
	Ol	·	·	·	·	·	<5	40	<5	20	20	n	n	n	n	n	50	60	50	30	10	n	n			
Cover of litter (%)																										
Number of species:																										
vascular plants		25	18	23	20	14	16	20	11	14	14	25	20	14	21	13	23	24	23	18	27	20	12			
mosses		1	1	1	2	3	7	3	6	4	4	9	3	4	8	3	1	6	8	3	1	5	2			
lichens		·	·	·	·	·	7	4	5	4	8	7	3	7	7	5	9	7	5	12	7	7	8			
1		2					3					4					5									
<i>Pinus sylvestris</i>	a	·	·	·	·	·	·	·	·	·	·	1	6	·	·	·	[1]	·	3	04	·	[04]	[04]			
	b	·	·	·	·	·	·	2	[04]	1	2	·	1	·	1	04	1	3	·	04	1	·	1			
<i>Populus tremula</i>	c	·	·	·	·	·	02	·	01	·	·	·	·	·	·	·	01	01	01	01	·	01	01			
	a	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
<i>Betula pendula</i>	b	·	·	·	·	·	02	·	04	·	1	04	04	04	04	·	·	·	·	·	3	3				
	c	·	·	·	·	·	·	·	04	·	·	·	02	04	·	·	·	·	·	·	·	1	04			
<i>Salix caprea</i>	a	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	[2]	·	·				
	b	·	·	·	·	·	·	1	·	·	01	04	01	04	04	04	·	·	04	2	·	·				
<i>Salix caprea</i>	c	·	·	·	·	·	·	·	·	·	·	·	01	·	·	01	01	·	·	01	·	·				
	a	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
<i>Picea abies</i>	b	·	·	·	·	·	3	·	·	[04]	01	·	·	·	·	·	·	·	·	·	·	·				
	c	·	·	·	·	·	·	·	·	·	·	·	·	·	·	01	·	·	·	·	·	·				
<i>Juniperus communis</i>	a	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
	b	01	·	·	·	·	·	·	1	1	04	04	04	04	01	·	04	1	04	·	1	04				
<i>Salix repens</i>	c	·	·	·	01	·	04	04	·	01	04	01	01	01	02	02	01	01	01	01	01	02	04			
	subsp. <i>rosmarinifolia</i>	b	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	04			
<i>Salix cinerea</i>	b	·	·	·	·	·	·	·	1	·	·	04	·	·	·	·	·	·	·	·	·	·				
	b	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
<i>Salix aurita</i>	b	·	·	·	·	·	·	·	·	·	·	01	·	·	·	·	·	·	·	·	·	·				
	b	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
<i>Frangula alnus</i>	c	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
	b	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
<i>Viburnum opulus</i>	b	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
	c	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
<i>Rosa</i> sp.	b	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
	c	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
<i>Rhamnus cathartica</i>	b	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
	c	02	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
<i>Pyrus communis</i>	c	·	·	·	·	·	·	·	·	·	·	01	01	·	·	·	·	·	·	·	·	·				
	c	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
<i>Quercus robur</i>	c	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
	c	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
<i>Padus serotina</i>	c	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
	c	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
<i>Sorbus aucuparia</i>	c	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
	c	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
Ch <i>Aperetalia</i>																										
<i>Vicia hirsuta</i>	02	1	1	04	01	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
	·	·	2	2	01	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·	·				
<i>Vicia angustifolia</i>																										



Succession stage	Initial																			
Succession phase	2					4														
Number of plot	82					67					84					61				
Ch <i>Agropyro-Rumicion crispi</i>																				
<i>Rumex crispus</i>	.	.	.	.	04	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
Ch <i>Artemisieta, Onopordion*</i>																				
<i>Artemisia vulgaris</i>	.	.	.	02	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
* <i>Berteroa incana</i>	.	.	01	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
* <i>Melandrium album</i>	02	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
Ch <i>Isoëto-Nanojuncetea, Cyperetalia fusci*, Radiolion linoidis**</i>																				
* <i>Gnaphalium uliginosum</i>	.	.	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
** <i>Hypericum humifusum</i>	.	.	.	.	.	.	01	.	.	.	.	.	.	.	.	01	01	01	01	01
Ch <i>Molinio-Arrhenatheretea, Molinietaia*, Arrhenatheretalia**, Cynosurion***</i>																				
<i>Poa pratensis</i>	2	2	2	3	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Vicia cracca</i>	01	04	02	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Rumex acetosa</i>	02	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Achillea millefolium</i>	04	1	1	2	2	1	01	1	02	01	02	01	02	02	.	01	01	01	01	04
<i>Holcus lanatus</i>	.	.	.	.	01	02	.	.	.	01	.	.	.	.	.	.	.	.	.	.
* <i>Juncus conglomeratus</i>	02	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
** <i>Taraxacum officinale</i> s.l.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
*** <i>Trifolium repens</i>	01	.	01	04	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
Ch <i>Koelerietea</i>																				
<i>Potentilla argentea</i>	01	01	01	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Brachythecium albicans</i>	.	1	1	1	01	04	04	.	.	.	04	.	.	02	.	.	.	01	.	.
<i>Hieracium pilosella</i>	01	01	04	2	.	1	5	4	02	2	1	04	04	04	.	04	02	02	01	2
<i>Rumex acetosella</i>	.	01	02	.	.	1	01	04	01	01	01	02	01	.	02	01	04	02	01	1
<i>Trifolium arvense</i>	.	02	.	.	.	.	01	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Rumex tenuifolius</i>	.	.	.	01	.	02	02	02	.	02	.	01	04	04	01	.	02	01	.	02
<i>Artemisia campestris</i>	01	.	.	.	.	.	2	.	.	.	.	02	.	04	.	01	01	.	.	01
<i>Scleranthus perennis</i>	.	.	.	.	.	01	.	.	.	01	02	04	01	04	02	01	01	02	01	02
<i>Jasione montana</i>	.	.	.	.	.	02	01	01	.	01	01	01	.	02	01	02	02	01	01	02
<i>Ceratodon purpureus</i>	.	.	.	.	.	1	.	1	.	1	01	.	.	01	.	.	.	04	04	.
<i>Cladonia furcata</i>	.	.	.	.	.	.	.	.	.	.	02	.	01	02	.	04	.	.	01	01
var. <i>palamea</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Veronica dillenii</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	.	.	.	.
<i>Festuca ovina</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	02	.	.	.	.	.	.
<i>Niphotrichum canescens</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
Ch <i>Corynephorretalia canescentis, Corynephorion canescentis*</i>																				
<i>Polytrichum piliferum</i>	.	.	.	.	.	04	4	5	4	7	1	1	04	1	1	4	2	2	4	2
<i>Corynephorus canescens</i>	.	.	.	.	.	01	01	1	.	04	04	04	02	02	04	02	1	01	04	02
<i>Cladonia arbuscula</i>	.	.	.	.	.	01	.	01	01	01	02	01	01	01	02	2	1	04	04	04
subsp. <i>mitis</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Cladonia cervicornis</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	01
<i>Cetraria aculeata</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Cladonia uncialis</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
* <i>Teesdalea nudicaulis</i>	.	.	.	.	.	.	.	.	01	.	.	.	.	.	.	01	01	.	.	01
* <i>Spergula morisonii</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
Ch <i>Festuco-Sedetalia, Armerion elongatae*, Koelerion glaucae**</i>																				
<i>Thymus serpyllum</i>	.	.	.	.	.	.	.	1	.	.	.	.	.	.	.	.	01	.	01	04
<i>Helichrysum arenarium</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	01	.	01
<i>Sedum maximum</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
* <i>Cerastium arvense</i>	.	.	02	1	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.





Succession stage	Initial																							
Succession phase	2				4												5							
Number of plot	82				67				84				61											
* <i>Armeria maritima</i> subsp. <i>elongata</i>	.	.	.	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
** <i>Gypsophila fastigiata</i>	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	.	.	.	.	.	.	.	.	.
** <i>Astragalus arenarius</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	.	.	.
** <i>Koeleria glauca</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
Ch <i>Trifolium medii</i>																								
<i>Galium mollugo</i>	04	1	.	04	.	.	.	.	.	02	.	.	.	.	.	.	.	.	.	.	.	.	.	.
Ch <i>Nardo-Callunetea</i>																								
<i>Calluna vulgaris</i>	.	.	.	.	.	02	.	.	.	2	.	.	04	2	04	6	04	4	02	02				
<i>Luzula campestris</i>	.	.	.	.	01	.	.	.	.	01	.	.	01	.	.	.	.	.	.	.	01			
<i>Danthonia decumbens</i>	.	.	.	.	.	.	.	.	.	01	.	.	.	.	04	.	.	.	.	.	.	.	.	.
<i>Luzula multiflora</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
Ch <i>Festuco-Brometea</i>																								
<i>Centaurea stoebe</i>	.	.	.	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Carlina vulgaris</i>	.	.	.	.	.	.	.	.	.	.	.	02	.	01	.	.	.	01	.	.	.	.	.	.
<i>Achillea collina</i>	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	01	.	.	.	.	.	.	.	.
Ch <i>Quercetea robori-petraeae</i>																								
<i>Hieracium sabaudum</i>	.	.	.	.	.	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Holcus mollis</i>	.	.	.	.	.	.	.	.	.	01	.	01	.	.	.	.	.	.	.	.	.	.	.	.
Ch <i>Dicrano-Pinion</i>																								
<i>Dicranum polysetum</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Monotropa hypopitys</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Cladonia gracilis</i>	.	.	.	.	.	.	.	.	.	.	.	.	1	1	.	04	04	.	02					
<i>Ptilidium ciliare</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Cladonia arbuscula</i> subsp. <i>arbuscula</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Chimaphila umbellata</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
Ch <i>Epilobietea, Epilobion angustifolii*</i>																								
<i>Calamagrostis epigejos</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	1	.	02	.	.	.	.	.	.	.	.
<i>Fragaria vesca</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
* <i>Chamaenerion angustifolium</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
Accompanying species																								
<i>Elymus repens</i>	02	04	.	4	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Vicia</i> sp.	1	2	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Cirsium arvense</i>	.	02	.	3	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Silene</i> sp.	01	02	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Viola arvensis</i>	.	01	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Carex hirta</i>	04	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Conyza canadensis</i>	.	.	.	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Lathyrus sylvestris</i>	.	.	.	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Oenothera</i> sp.	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Viola tricolor</i>	.	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Stellaria graminea</i>	3	3	3	2	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Solidago virgaurea</i>	04	01	1	1	1	1	2	2	04	04	04	2	1	2	2	04	04	1	2	1				
<i>Knautia arvensis</i>	01	01	04	02	02	04	.	04	.	01	01	.	.	01	.	01	.	01	.	.	.	.	.	.
<i>Convolvulus arvensis</i>	02	01	01	02	04	01	.	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Hypericum perforatum</i>	01	01	.	01	02	.	01	.	01	.	.	.	01	01	02	.	04	02	.					
<i>Agrostis capillaris</i>	2	2	1	.	6	5	01	01	.	04	.	01	01	01	01	01	01	01	01	.	.	.	.	.
<i>Sciuro-hypnum oedipodium</i>	.	.	02	02	01	01	.	01	.	01	.	01	.	01	.	01	.	.	.	.	.	.	.	.
<i>Polytrichum commune</i>	2	.	.	.	04	02	1	04	1	2	5	6	2	.	.	.	.	.	.	.	.	.	.	.

[illegible]

Succession stage	Initial																						
Succession phase	2					4											5						
Number of plot	82					67					84						61						
<i>Agrostis gigantea</i>	.	.	.	.	2	.	.	.	.	.	.	.	.	.	.	.	.	.	01	02	.	.	
<i>Pleurozium schreberi</i>	.	.	.	.	1	02	.	.	.	.	04	.	04	01	.	.	.	02	02	.	.	01	.
<i>Peucedanum oreoselinum</i>	.	.	.	1	.	.	.	.	.	.	.	02	.	.	.	.	.	02	.	.	.	02	.
<i>Festuca rubra</i>	.	.	04	.	.	.	.	.	.	.	01	.	01	.	.	.	01	01	01	01	04	.	.
<i>Euphrasia stricta</i>	.	.	.	01	.	.	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Polytrichum juniperinum</i>	.	.	.	.	.	1	.	01	3	1	4	3	2	1	1	.	1	.	04	.	2	3	
<i>Melampyrum pratense</i>	.	.	.	.	.	.	.	.	.	.	04	1	04	04	02	.	.	.	.	.	04	04	
<i>Poa angustifolia</i>	.	.	.	.	.	.	1	.	01	.	04	.	02	01	.	.	.	.	.	.	.	02	
<i>Cladonia chlorophaea</i> s.l.	.	.	.	.	.	02	04	04	1	02	01	01	.	01	.	.	.	02	02	.	02	.	
<i>Cladonia cornuta</i>	.	.	.	.	.	01	.	02	01	1	04	.	04	01	04	1	2	.	04	1	.	1	
<i>Cladonia</i> cf. <i>glauca</i>	.	.	.	.	.	2	1	2	2	01	.	.	.	01	.	.	.	.	.	.	.	02	
<i>Cladonia phyllophora</i>	.	.	.	.	.	01	.	.	01	02	.	02	.	02	1	1	04	1	1	04	04		
<i>Cladonia furcata</i> var. <i>furcata</i>	.	.	.	.	.	.	.	.	.	01	04	04	02	.	.	02	.	01	01	01	04	04	
<i>Cladonia subulata</i>	.	.	.	.	.	02	01	.	.	.	.	01	01	.	.	.	.	01	.	.	.	.	
<i>Hypochoeris radicata</i>	.	.	.	.	.	01	.	.	.	.	01	01	.	.	.	01	.	01	.	.	01	.	
<i>Cladonia furcata</i> f. <i>fissa</i>	.	.	.	.	.	.	.	.	.	.	.	.	01	01	01	.	.	.	.	.	.	.	
<i>Placynthiella uliginosa</i>	.	.	.	.	.	04	.	.	.	02	.	.	.	.	.	02	02	.	04	.	01	.	
<i>Hypnum cupressiforme</i>	.	.	.	.	.	.	.	.	04	.	02	.	.	.	.	.	02	04	.	.	02	.	
<i>Carex ericetorum</i>	.	.	.	.	.	.	.	.	02	.	.	.	.	02	.	.	.	.	.	.	.	.	
<i>Euphrasia</i> sp.	.	.	.	.	.	.	.	.	.	.	02	.	.	01	.	.	.	.	.	.	02	.	
<i>Cladonia fimbriata</i>	.	.	.	.	.	01	.	.	.	.	01	.	.	.	.	.	.	.	.	.	01	02	
<i>Carex ovalis</i>	.	.	.	.	.	02	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	
<i>Hieracium umbellatum</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	.	.	.	01	02	.	
<i>Cephaloziella starkei</i>	.	.	.	.	.	.	.	.	.	.	01	.	.	.	.	.	02	.	.	.	01	.	
<i>Brachythecium</i> <i>salebrosum</i>	.	.	.	.	.	.	.	01	.	.	.	.	.	.	.	.	.	04	.	.	.	.	
<i>Brachytheciastrium</i> <i>velutinum</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	.	01	.	.	.	.	
<i>Pohlia nutans</i>	.	.	.	.	.	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	
<i>Cladonia floerkeana</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	.	01	.	.	.	
<i>Cladonia scabriuscula</i>	.	.	.	.	.	01	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	
<i>Cladonia pleurota</i>	.	.	.	.	.	.	.	.	.	01	.	.	.	.	.	.	.	.	.	.	.	.	
<i>Cladonia crispata</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	01	.	02	
<i>Euphrasia vernalis</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	01	.	.	01	.	.	
<i>Chamaecytisus</i> <i>ratisbonensis</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	1	.	2	.	.	
<i>Linaria vulgaris</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	01	.	.	
<i>Erigeron acris</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	
<i>Viola canina</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	
<i>Viola</i> sp. 1	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	
<i>Silene vulgaris</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	.	.	.	
<i>Trapeliopsis granulosa</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	.	
<i>Cladonia coccifera</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	.	.	.	
<i>Stereocaulon</i> <i>condensatum</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	.	
<i>Cladonia deformis</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	01	.	.	.	.	.	.	
<i>Cladonia rangiferina</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	
<i>Pteridium aquilinum</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	
<i>Polytrichiastrum</i> <i>formosum</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	
<i>Cerastium holosteoides</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	

[illegible]

Succession stage	Initial																							
Succession phase	2					4											5							
Number of plot	82					67					84						61							
<i>Luzula</i> sp.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Amblystegnum serpens</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Cladonia macilenta</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Cladonia squamosa</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Cetraria islandica</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Vincetoxicum</i> <i>hirundinaria</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Dicranum scoparium</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Lophocolea bidentata</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Lophocolea heterophylla</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Pseudoscleropodium</i> <i>purum</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Plagiomnium affine</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Dicranum</i> sp.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Placynthiella oligotropha</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Cladonia coniocraea</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Pimpinella saxifraga</i>	.	04	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Genista tinctoria</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Viola</i> sp. 2	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Veronica officinalis</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Viola</i> sp. 3	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Saxifraga</i> sp.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Galium verum</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Veronica chamaedrys</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Carex spicata</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Erophila verna</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Luzula pallescens</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Thuidium tamariscinum</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.
<i>Plagiomnium cuspidatum</i>	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.

[illegible]



Table A2.2  
Vegetation description in the permanent plots in the Czechy Orlańskie reserve  
(based on FALIŃSKI 1992, changed; only the plots used for mycological examination are included;  
Braun-Blanquet method)

Succession phase		9/10				
Number of plot		2C	5C	6C	7C	8C
Relevé area (m <sup>2</sup> )		200	200	200	200	200
Cover of plant layers (%)						
	a	60	60	50	60	60
	a <sub>1</sub>	60	60	50	60	60
	a <sub>2</sub>	10	10	·	10	·
	b	10	30	20	10	+
	c	90	80	70	80	70
	d	100	90	100	100	100
Number of species		28	26	31	24	22
1		2	3	4	5	6
<i>Pinus sylvestris</i> <sup>1</sup>	a	4.4	4.4	4.4	4.4	4.4
	c	·	·	+	+	·
<i>Picea abies</i> <sup>2</sup>	a	1.1	2.1	·	·	·
	b	1.2	1.1	·	1.2	+
	c	+	+	·	+	+
<i>Betula pendula</i> <sup>2</sup>	a	·	·	·	1.1	·
	b	+ .2	1.2	1.1	·	+
	c	+	+	·	·	+
<i>Betula pubescens</i> <sup>2</sup>	a	·	2.2	·	·	·
	c	+	·	·	·	·
<i>Quercus petraea</i> <sup>2</sup>	a	·	2.1	·	·	·
	c	·	·	·	·	+
<i>Salix caprea</i>	b	1.1	·	·	·	·
<i>Frangula alnus</i>	b	1.1	2.2	2.2	1.1	1.1
	c	+	+	+	+	·
<i>Juniperus communis</i>	b	·	3.3	2.3	1.2	+
	c	+	+	+	+	+
<i>Populus tremula</i>	c	+	·	·	·	·
<i>Ch Peucedano-Pinetum, Dicrano-Pinion</i>						
<i>Chamaecytisus ratisbonensis</i>		1.2	1.2	2.2	2.2	1.2
<i>Chimaphila umbellata</i>		+	+ .2	1.2	+	+ .2
<i>Pyrola chlorantha</i>		+	·	+	·	+
<i>Diphasiastrum complanatum</i>		·	·	2.3	·	·
<i>Ch Quercu-Piceetum, Vaccinio-Piceion</i>						
<i>Goodyera repens</i>		·	·	·	·	+
<i>Ch Vaccinio-Piceetalia, Vaccinio-Piceetea</i>						
<i>Vaccinium myrtillus</i>		2.3	4.4	2.2	4.4	3.4
<i>Vaccinium vitis-idaea</i>		4.4	2.2	3.4	3.3	3.3
<i>Dicranum polysetum</i>		+ .2	1.2	1.2	1.2	1.2
<i>Ptilium crista-castrensis</i>		+	+ .2	+ .2	+ .2	+ .2
<i>Melampyrum pratense</i>		+ .2	+	1.2	1.2	+
<i>Orthilia secunda</i>		·	+ .2	+ .2	·	+ .2
<i>Trientalis europaea</i>		+	·	+	·	·
Accompanying species						
<i>Luzula pilosa</i>		1.2	+ .2	+ .2	+	+

<i>Pleurozium schreberi</i>		3.3	4.4	3.4	3.4	4.4
<i>Hylocomium splendens</i>		4.4	3.4	3.4	3.4	3.4
<i>Polytrichum commune</i>		+2	+2	+2	+2	+2
<i>Dryopteris carthusiana</i>		.	+2	+	.	.
<i>Festuca rubra</i>		+2	+	+	+2	.
<i>Festuca ovina</i>		1.2	2.2	1.2	1.2	1.2
<i>Rubus idaeus</i>	b/c	.	.	+	.	.
<i>Lycopodium clavatum</i>		.	1.2	.	.	2.2
<i>Pteridium aquilinum</i>		.	1.2	.	.	.
<i>Hieracium lachenalii</i>		.	+	+	+	.
<i>Agrostis capillaris</i>		.	+	+	+	.
<i>Rubus saxatilis</i>		+	.	.	.	.
<i>Acer platanoides</i>		+	.	+	+	.
<i>Polytrichum juniperinum</i>		.	.	.	+2	+2
<i>Solidago virgaurea</i>		.	.	+	.	.
<i>Carpinus betulus</i>	c	+	.	.	+	.
<i>Hieracium pilosella</i>		.	.	.	+	.
<i>Rumex acetosella</i>		+2	.	.	.	.

Sporadic species: plot 2C – *Veronica officinalis* +; plot 6C – *Danthonia decumbens* +, *Carex ericetorum* 1.2, *Agrostis alba* [= *A. gigantea* or *A. stolonifera*] +, *Hieracium umbellatum* +; plot 7C – *Carex ovalis* +2

<sup>1</sup> layer a refers to layer a<sub>1</sub> in FALIŃSKI (1992); <sup>2</sup> layer a refers to layer a<sub>2</sub> in FALIŃSKI (1992)



## Appendix 3

### Macrofungi in the plots representing subsequent phases of vegetation succession in the old-fields near the Białowieża old-growth forest

Table A3.1  
Macrofungi in plot 82 (phase 2, Jelonka reserve)

Species	AC '95	AC '96	AC2y	aACy	mACv	SF%	tTF	TFy
<i>Crinipellis scabellia</i>	43	412	455	227.5	366	100	6	2
<i>Marasmius oreades</i>	103	123	226	113.0	61	100	10	2
<i>Mycena flavoalba</i>	157	52	209	104.5	157	100	2	2
<i>Mycena olivaceomarginata</i>	96	16	112	56.0	76	90	4	2
<i>Mycena leptcephala</i>	164	·	164	82.0	164	80	1	1
<i>Galerina</i> spp.	40	10	50	25.0	39	70	3	2
<i>Coprinopsis friesii</i>	·	33	33	16.5	31	60	2	1
<i>Mycena pura</i>	1	5	6	3.0	2	40	4	2
<i>Parasola plicatilis</i>	4	6	10	5.0	6	30	2	2
<i>Cystoderma amianthinum</i>	5	4	9	4.5	5	10	2	2
<i>Clitocybe diatreta</i>	8	·	8	4.0	8	10	1	1
<i>Rickenella fibula</i>	6	·	6	3.0	6	10	1	1
<i>Clitocybe brumalis</i>	1	·	1	0.5	1	10	1	1
<i>Conocybe spiculoides</i>	1	·	1	0.5	1	10	1	1
<i>Entoloma undatum</i>	1	·	1	0.5	1	10	1	1

AC – abundance of carpophores during the indicated year; AC2y – abundance of carpophores during 2 years of observations; aACy – average density of carpophores per year; mACv – maximum density of carpophores per visit; SF% – spatial frequency of a species (percentage of subplots with a species presence; plot surface – 1000 m<sup>2</sup>, number of subplots – 10); tTF – number of visits with a species presence; TFy – number of years with a species presence

Table A3.2  
Macrofungi in plot 67 (phase 4, Jelonka reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Inocybe lacera</i>	1146	1107	216	2469	823.0	865	92.5	9	3
<i>Suillus luteus</i>	4	78	140	222	74.0	72	85.0	8	3
<i>Amanita muscaria</i>	7	124	134	265	88.3	127	65.0	6	3
<i>Chalciporus piperatus</i>	·	49	57	106	35.3	28	57.5	5	2
<i>Laccaria proxima</i>	9	67	93	169	56.3	67	42.5	5	3
<i>Hygrophorus hypothejus</i>	11	14	37	62	20.7	22	37.5	4	3
<i>Cortinarius</i> spp. I	2	167	110	279	93.0	156	35.0	5	3
<i>Cortinarius cinnamomeoluteus</i>	4	17	35	56	18.7	31	25.0	6	3
<i>Inocybe asterospora</i>	·	20	·	20	6.7	20	15.0	1	1
<i>Cortinarius</i> spp. II	·	·	17	17	5.7	10	15.0	2	2
<i>Cortinarius</i> spp. III	·	18	3	21	7.0	18	12.5	2	2
<i>Hebeloma</i> sp. 2	·	8	1	9	3.0	8	12.5	2	2
<i>Cortinarius purpurascens</i>	·	3	11	14	4.7	10	10.0	3	2
<i>Leccinum aurantiacum</i>	·	·	5	5	1.7	4	10.0	2	1

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Hebeloma</i> sp. 1	·	4	1	5	1.7	2	7.5	3	2
<i>Boletus edulis</i>	·	·	4	4	1.3	3.0	7.5	2	1
<i>Paxillus involutus</i>	3	3	·	6	2.0	3	5.0	2	2
<i>Xerocomus subtomentosus</i>	·	5	·	5	1.7	5	5.0	1	1
<i>Clitopilus prunulus</i>	·	3	·	3	1.0	2	5.0	2	1
<i>Inocybe calida</i>	·	4	·	4	1.3	4	2.5	1	1
<i>Hygrophorus aureus</i>	·	4	·	4	1.3	4	2.5	1	1
<i>Thelephora terrestris</i>	3	·	·	3	1.0	3	2.5	1	1
<i>Rhodocollybia butyracea</i> f. <i>asema</i>	·	·	3	3	1.0	3	2.5	1	1
<i>Ramaria eumorpha</i>	·	·	3	3	1.0	3	2.5	1	1
<i>Russula fragilis</i>	·	1	·	1	0.3	1	2.5	1	1
<b>Other species</b>									
<i>Psilocybe montana</i>	208	183	152	543	181.0	192	82.5	10	3
<i>Cantharellula umbonata</i>	14	571	735	1320	440.0	362	72.5	9	3
<i>Entoloma</i> spp.	19	56	34	109	36.3	56	70.0	5	3
<i>Mycena olivaceomarginata</i>	44	13	·	57	19.0	26	62.5	3	2
<i>Rickenella fibula</i>	6	94	1	101	33.7	52	55.0	5	3
<i>Clitocybe diatreta</i>	21	80	12	113	37.7	71	52.5	9	3
<i>Clavaria argillacea</i>	167	83	94	344	114.7	167	50.0	5	3
<i>Cystoderma amianthinum</i>	3	48	27	78	26.0	26	42.5	7	3
<i>Galerina atkinsoniana</i>	1	140	19	160	53.3	75	37.5	6	3
<i>Mycetinis scorodoni</i>	42	67	141	250	83.3	129	32.5	8	3
<i>Collybia cirrhata</i>	102	20	8	130	43.3	88	32.5	4	3
<i>Mycena flavoalba</i>	7	142	25	174	58.0	130	30.0	4	3
<i>Mycena leptcephala</i>	1	42	·	43	14.3	42	27.5	2	2
<i>Psathyrella gyroflexa</i>	·	19	·	19	6.3	11	27.5	2	1
<i>Galerina</i> spp.	2	10	14	26	8.7	12	22.5	5	3
<i>Lycoperdon molle</i>	·	10	24	34	11.3	16	12.5	5	2
<i>Panaeolus papilionaceus</i>	·	28	2	30	10.0	17	12.5	3	2
<i>Gymnopus androsaceus</i>	15	·	8	23	7.7	15	10.0	3	2
<i>Lycoperdon nigrescens</i>	·	2	5	7	2.3	4	10.0	3	2
<i>Ampulloclitocybe clavipes</i>	·	9	1	10	3.3	9	7.5	2	2
<i>Lycoperdon perlatum</i>	·	5	·	5	1.7	5	5.0	1	1
<i>Gymnopus dryophilus</i>	4	·	·	4	1.3	4	5.0	1	1
<i>Hygroaster asterosporus</i>	·	2	1	3	1.0	2	5.0	2	2
<i>Lycoperdon dermoxanthum</i>	·	1	1	2	0.7	1	5.0	2	2
<i>Rhodocybe caelata</i>	2	·	·	2	0.7	1	5.0	2	1
<i>Galerina hypnorum</i>	·	5	·	5	1.7	5	2.5	1	1
<i>Mycena vitilis</i>	·	·	2	2	0.7	2	2.5	1	1
<i>Galerina pumila</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Mycena pura</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Rhodocybe hirneola</i>	1	·	·	1	0.3	1	2.5	1	1

AC – abundance of carpophores during the indicated year; AC3y – abundance of carpophores during 3 years of observations; aDCy – average density of carpophores per year; mDCv – maximum density of carpophores per visit; SF% – spatial frequency of a species (percentage of subplots with the species presence; plot surface – 1000 m<sup>2</sup>, number of subplots – 40); tTF – number of visits with a species presence; TFy – number of years with a species presence; x – perennial carpophores

Table A3.3  
Macrofungi in plot 84 (phase 4, Jelonka reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Inocybe lacera</i>	431	191	139	761	253.7	216	82.5	6	3
<i>Cortinarius cinnamomeoluteus</i>	62	393	316	771	257.0	365	75.0	9	3
<i>Amanita muscaria</i>	4	98	91	193	64.3	98	75.0	5	3
<i>Laccaria proxima</i>	88	253	106	447	149.0	205	70.0	8	3
<i>Cortinarius</i> spp. I	10	147	55	212	70.7	146	67.5	5	3
<i>Suillus luteus</i>	5	20	51	76	25.3	46	57.5	6	3
<i>Hygrophorus hypothejus</i>	17	27	37	81	27.0	34	45.0	5	3
<i>Russula fragilis</i>	1	61	3	65	21.7	61	37.5	4	3
<i>Leccinum aurantiacum</i>	1	4	9	14	4.7	7	30.0	4	3
<i>Cortinarius</i> spp. II	·	58	11	69	23.0	58	22.5	2	2
<i>Amanita pantherina</i>	5	2	2	9	3.0	5	17.5	3	3
<i>Thelephora terrestris</i>	3	20	22	45	15.0	20	15.0	6	3
<i>Chalciporus piperatus</i>	·	6	6	12	4.0	4	15.0	4	2
<i>Inocybe asterospora</i>	·	35	·	35	11.7	35	12.5	1	1
<i>Laccaria bicolor</i>	1	14	8	23	7.7	14	10.0	3	3
<i>Russula galochroa</i>	·	4	7	11	3.7	7	10.0	3	2
<i>Thelephora caryophyllea</i>	6	·	·	6	2.0	5	10.0	2	1
<i>Cortinarius</i> spp. III	·	32	·	32	10.7	32	7.5	1	1
<i>Amanita rubescens</i>	·	7	4	11	3.7	6	7.5	3	2
<i>Clitopilus prunulus</i>	·	8	·	8	2.7	8	7.5	1	1
<i>Boletus edulis</i>	·	3	3	6	2.0	3	7.5	2	2
<i>Tricholoma portentosum</i>	·	11	3	14	4.7	11	5.0	2	2
<i>Inocybe calida</i>	·	6	·	6	2.0	6	5.0	1	1
<i>Paxillus involutus</i>	·	3	·	3	1.0	3	5.0	1	1
<i>Amanita citrina</i>	1	2	·	3	1.0	2	5.0	2	2
<i>Gyroporus castaneus</i>	·	1	2	3	1.0	2	5.0	2	2
<i>Coltricia perennis</i>	1	1	1	3	1.0	1	5.0	3	3
<i>Hebeloma</i> sp. 1	·	2	·	2	0.7	2	5.0	1	1
<i>Leccinum scabrum</i>	1	1	·	2	0.7	1	5.0	2	2
<i>Russula galochroa</i>	·	2	·	2	0.7	2	5.0	1	1
<i>Tricholoma albobrunneum</i>	1	1	·	2	0.7	1	5.0	2	2
<i>Tricholoma saponaceum</i> var. <i>squamosum</i>	·	25	·	25	8.3	25	2.5	1	1
<i>Inocybe praetervisa</i>	·	3	·	3	1.0	3	2.5	1	1
<i>Russula emetica</i>	·	3	·	3	1.0	3	2.5	1	1
<i>Boletus reticulatus</i>	·	·	2	2	0.7	2	2.5	1	1
<i>Inocybe assimilata</i>	·	2	·	2	0.7	2	2.5	1	1
<i>Lactarius vietus</i>	·	2	·	2	0.7	2	2.5	1	1
<i>Russula subfoetens</i>	·	2	·	2	0.7	2	2.5	1	1
<i>Xerocomus subtomentosus</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Rhodocollybia butyracea</i> f. <i>asema</i>	·	·	1	1	0.3	1	2.5	1	1
<b>Other species</b>									
<i>Clitocybe diatreta</i>	103	187	39	329	109.7	124	92.5	8	3
<i>Psilocybe montana</i>	52	106	37	195	65.0	50	85.0	11	3
<i>Entoloma</i> spp.	37	262	4	303	101.0	223	82.5	5	3
<i>Clavaria argillacea</i>	428	180	483	1091	363.7	428	70.0	5	3
<i>Cystoderma amianthinum</i>	8	80	107	195	65.0	63	57.5	8	3
<i>Galerina</i> spp.	12	201	·	213	71.0	200	55.0	4	2
<i>Rhodocybe parilis</i>	18	21	2	41	13.7	15	40.0	6	3
<i>Lycoperdon dermoxanthum</i>	4	16	3	23	7.7	15	32.5	4	3

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Rickenella fibula</i>	3	16	1	20	6.7	14	30.0	5	3
<i>Gymnopus androsaceus</i>	157	145	·	302	100.7	157	22.5	4	2
<i>Collybia cirrhata</i>	3	36	10	49	16.3	30	22.5	5	3
<i>Mycena olivaceomarginata</i>	6	7	·	13	4.3	6	20.0	3	2
<i>Lycoperdon perlatum</i>	5	13	2	20	6.7	6	17.5	6	3
<i>Clitocybe vibecina</i>	32	·	3	35	11.7	32	15.0	2	2
<i>Entoloma jubatum</i>	·	·	13	13	4.3	13	15.0	1	1
<i>Mycena leptcephala</i>	·	26	·	26	8.7	26	12.5	1	1
<i>Mycena pura</i>	4	2	·	6	2.0	4	12.5	2	2
<i>Galerina atkinsoniana</i>	4	20	10	34	11.3	14	10.0	5	3
<i>Mycetinis scorodonius</i>	9	14	·	23	7.7	14	10.0	2	2
<i>Mycena flavaalba</i>	·	10	·	10	3.3	10	10.0	1	1
<i>Ampulloclitocybe clavipes</i>	·	5	1	6	2.0	5	10.0	2	2
<i>Flammulaster granulosus</i>	11	6	·	17	5.7	11	7.5	2	2
<i>Hygroaster asterosporus</i>	·	5	·	5	1.7	5	7.5	1	1
<i>Galerina hypnorum</i>	1	2	1	4	1.3	2	7.5	3	3
<i>Macrolepiota procera</i>	1	2	1	4	1.3	2	7.5	3	3
<i>Lycoperdon nigrescens</i>	1	2	·	3	1.0	2	7.5	2	2
<i>Galerina pumila</i>	·	15	12	27	9.0	15	5.0	2	2
<i>Hygrocybe virginea</i> var. <i>virginea</i>	·	2	12	14	4.7	12	5.0	2	2
<i>Entoloma rhodopolium</i>	·	2	·	2	0.7	2	5.0	1	1
<i>Mycena metata</i>	·	2	·	2	0.7	2	5.0	1	1
<i>Mycena stylobates</i>	·	1	1	2	0.7	1	5.0	2	2
<i>Dacrymyces stillatus</i>	·	50	·	50	16.7	50	2.5	1	1
<i>Crepidotus variabilis</i>	·	35	·	35	11.7	30	2.5	2	1
<i>Hypholoma fasciculare</i>	·	12	·	12	4.0	12	2.5	1	1
<i>Sphaerobolus stellatus</i>	·	5	·	5	1.7	5	2.5	1	1
<i>Tubaria furfuracea</i>	·	3	·	3	1.0	2	2.5	2	1
<i>Macrolepiota mastoidea</i>	2	·	·	2	0.7	2	2.5	1	1
<i>Conocybe rickeniana</i>	1	·	·	1	0.3	1	2.5	1	1
<i>Gymnopus dryophilus</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Gymnopus putillus</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Lycoperdon molle</i>	1	·	·	1	0.3	1	2.5	1	1
<i>Mycena aetites</i>	·	1	·	1	0.3	1	2.5	1	1

Abbreviations: see Tab. A3.2

Table A3.4  
Macrofungi in plot 61 (phase 5, Jelonka reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Suillus luteus</i>	5	202	120	327	109.0	130	92.5	8	3
<i>Cortinarius cinnamomeoluteus</i>	200	885	303	1388	462.7	846	90.0	11	3
<i>Hygrophorus hypothejus</i>	201	154	96	451	150.3	201	80.0	5	3
<i>Cortinarius</i> spp. I	49	206	93	348	116.0	165	60.0	8	3
<i>Amanita muscaria</i>	2	77	112	191	63.7	109	60.0	4	3
<i>Tricholoma albobrunneum</i>	3	63	51	117	39.0	53	42.5	5	3
<i>Laccaria proxima</i>	43	152	119	314	104.7	131	40.0	6	3
<i>Chalciporus piperatus</i>	2	19	9	30	10.0	13	35.0	6	3
<i>Cortinarius</i> spp. III	·	59	89	148	49.3	89	25.0	2	2
<i>Cortinarius</i> spp. II	·	20	9	29	9.7	20	20.0	3	2
<i>Rhodocollybia butyracea</i> f. <i>asema</i>	·	·	25	25	8.3	25	17.5	1	1
<i>Tricholoma portentosum</i>	5	9	7	21	7.0	9	17.5	4	3



Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Tricholoma saponaceum</i> var. <i>saponaceum</i>	3	·	29	32	10.7	23	15.0	3	2
<i>Inocybe lacera</i>	6	4	15	25	8.3	10	15.0	6	3
<i>Tricholoma equestre</i>	·	13	3	16	5.3	10	15.0	5	2
<i>Lactarius quieticolor</i>	·	16	1	17	5.7	15	7.5	3	2
<i>Thelephora terrestris</i>	·	2	5	7	2.3	3	7.5	3	2
<i>Cortinarius cinnabarinus</i>	·	1	3	4	1.3	3	7.5	2	2
<i>Cortinarius anthracinus</i>	·	3	·	3	1.0	3	7.5	1	1
<i>Paxillus involutus</i>	·	·	9	9	3.0	7	5.0	3	1
<i>Boletus edulis</i>	2	1	2	5	1.7	2	5.0	3	3
<i>Inocybe praetervisa</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Xerocomus subtomentosus</i>	·	·	1	1	0.3	1	2.5	1	1
<b>Other species</b>									
<i>Psilocybe montana</i>	162	81	64	307	102.3	82	82.5	12	3
<i>Clitocybe diatreta</i>	199	264	153	616	205.3	174	67.5	10	3
<i>Clavaria argillacea</i>	52	364	231	647	215.7	264	57.5	5	3
<i>Entoloma</i> spp.	18	71	2	91	30.3	41	55.0	6	3
<i>Collybia cirrhata</i>	127	322	7	456	152.0	322	35.0	4	3
<i>Rickenella fibula</i>	1	50	1	52	17.3	37	32.5	5	3
<i>Lycoperdon dermoxanthum</i>	1	14	3	18	6.0	13	30.0	4	3
<i>Mycena leptcephala</i>	73	14	·	87	29.0	73	27.5	2	2
<i>Mycena flavoalba</i>	4	62	3	69	23.0	54	17.5	4	3
<i>Mycena olivaceomarginata</i>	6	5	·	11	3.7	6	12.5	3	2
<i>Galerina atkinsoniana</i>	3	28	6	37	12.3	17	10.0	4	3
<i>Gymnopus dryophilus</i>	31	1	4	36	12.0	31	10.0	3	3
<i>Lycoperdon marginatum</i>	2	7	5	14	4.7	4	10.0	6	3
<i>Galerina</i> spp.	9	3	·	12	4.0	9	10.0	3	2
<i>Mycena pura</i>	1	3	1	5	1.7	3	10.0	3	3
<i>Gymnopus androsaceus</i>	59	·	·	59	19.7	59	7.5	1	1
<i>Galerina pumila</i>	·	16	4	20	6.7	13	7.5	4	2
<i>Entoloma jubatum</i>	·	5	5	10	3.3	5	7.5	2	2
<i>Macrolepiota procera</i>	3	3	·	6	2.0	3	7.5	2	2
<i>Psathyrella gyroflexa</i>	·	4	·	4	1.3	3	7.5	2	1
<i>Clitocybe fragrans</i>	·	·	31	31	10.3	31	5.0	1	1
<i>Collybia cookei</i>	2	20	5	27	9.0	20	5.0	3	3
<i>Hygroaster asterosporus</i>	4	1	·	5	1.7	4	5.0	2	2
<i>Mycena stylobates</i>	1	2	·	3	1.0	1	5.0	3	2
<i>Rhodocybe parilis</i>	1	1	·	2	0.7	1	5.0	2	2
<i>Hygrocybe virginea</i> var. <i>virginea</i>	11	14	11	36	12.0	14	2.5	3	3
<i>Cystoderma amianthinum</i>	·	17	·	17	5.7	17	2.5	1	1
<i>Mycena adonis</i>	2	15	·	17	5.7	15	2.5	2	2
<i>Crepidotus variabilis</i>	·	7	·	7	2.3	7	2.5	1	1
<i>Ampulloclitocybe clavipes</i>	3	·	·	3	1.0	3	2.5	1	1
<i>Gymnopus putillus</i>	3	·	·	3	1.0	3	2.5	1	1
<i>Leptosporomyces galzinii</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Macrolepiota mastoidea</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Marasmius oreades</i>	1	·	·	1	0.3	1	2.5	1	1
<i>Tapinella atrotomentosa</i>	·	·	1	1	0.3	1	2.5	1	1

Abbreviations: see Tab. A3.2

Table A3.5  
Macrofungi in plot 85 (phase 5, Jelonka reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Inocybe lacera</i>	584	337	229	1150	383.3	393	85.0	9	3
<i>Laccaria proxima</i>	159	214	208	581	193.7	161	75.0	8	3
<i>Suillus luteus</i>	5	53	107	165	55.0	64	70.0	8	3
<i>Hygrophorus hypothejus</i>	23	124	98	245	81.7	122	67.5	5	3
<i>Amanita muscaria</i>	16	78	138	232	77.3	134	67.5	7	3
<i>Cortinarius cinnamomeoluteus</i>	9	144	149	302	100.7	141	62.5	7	3
<i>Thelephora terrestris</i>	13	44	14	71	23.7	34	37.5	6	3
<i>Cortinarius</i> spp. I	1	68	32	101	33.7	56	35.0	5	3
<i>Chalciporus piperatus</i>	4	12	9	25	8.3	10	32.5	6	3
<i>Russula fragilis</i>	·	38	1	39	13.0	38	30.0	2	2
<i>Leccinum aurantiacum</i>	1	6	15	22	7.3	15	25.0	4	3
<i>Clitopilus prunulus</i>	13	8	1	22	7.3	13	15.0	4	3
<i>Amanita rubescens</i>	7	6	10	23	7.7	7	12.5	5	3
<i>Paxillus involutus</i>	·	3	16	19	6.3	16	10.0	2	2
<i>Inocybe curvipes</i>	·	·	12	12	4.0	10	10.0	2	1
<i>Cortinarius</i> spp. II	·	10	4	14	4.7	10	12.5	2	2
<i>Boletus edulis</i>	·	·	13	13	4.3	13	12.5	1	1
<i>Cortinarius</i> spp. III	·	4	·	4	1.3	4	7.5	1	1
<i>Xerocomus subtomentosus</i>	1	2	1	4	1.3	2	7.5	3	3
<i>Scleroderma fuscum</i>	·	3	·	3	1.0	3	7.5	1	1
<i>Gyroporus castaneus</i>	·	1	2	3	1.0	2	5.0	2	2
<i>Inocybe calida</i>	·	3	·	3	1.0	3	5.0	1	1
<i>Inocybe praetervisa</i>	·	1	2	3	1.0	2	5.0	2	2
<i>Amanita pantherina</i>	·	2	·	2	0.7	2	5.0	1	1
<i>Tricholoma albobrunneum</i>	·	·	2	2	0.7	2	5.0	1	1
<i>Tricholoma equestre</i>	·	3	·	3	1.0	3	2.5	1	1
<i>Laccaria bicolor</i>	·	2	·	2	0.7	2	2.5	1	1
<i>Xerocomus chrysenteron</i>	·	2	·	2	0.7	2	2.5	1	1
<i>Lactarius uvidus</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Rhodocollybia butyracea</i> f. <i>asema</i>	1	·	·	1	0.3	1	2.5	1	1
<i>Russula clavipes</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Russula galochroa</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Russula subfetens</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Russula vinosobrunnea</i>	·	1	·	1	0.3	1	2.5	1	1
<b>Other species</b>									
<i>Psilocybe montana</i>	75	217	107	399	133.0	97	97.5	13	3
<i>Clitocybe diatreta</i>	119	127	41	287	95.7	72	65.0	9	3
<i>Entoloma</i> spp.	47	84	3	134	44.7	48	57.5	6	3
<i>Galerina</i> spp.	10	43	56	109	36.3	53	42.5	6	3
<i>Clavaria argillacea</i>	64	141	306	511	170.3	168	35.0	5	3
<i>Rickenella fibula</i>	3	83	8	94	31.3	73	35.0	6	3
<i>Cystoderma amianthinum</i>	2	34	52	88	29.3	32	35.0	7	3
<i>Rhodocybe parilis</i>	15	32	·	47	15.7	24	35.0	4	2
<i>Lycoperdon dermoxanthum</i>	7	15	·	22	7.3	14	27.5	3	2
<i>Mycena leptcephala</i>	2	18	·	20	6.7	18	22.5	2	2
<i>Entoloma jubatum</i>	1	4	12	17	5.7	12	22.5	3	3
<i>Cantharellula umbonata</i>	3	46	105	154	51.3	55	17.5	7	3
<i>Collybia cirrhata</i>	10	52	·	62	20.7	52	17.5	3	2
<i>Crinipellis scabellula</i>	26	6	·	32	10.7	21	17.5	3	2
<i>Lycoperdon perlatum</i>	1	11	9	21	7.0	6	15.0	5	3

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Mycena flavoalba</i>	1	18	1	20	6.7	18	15.0	3	3
<i>Galerina hypnorum</i>	·	22	31	53	17.7	31	12.5	3	2
<i>Lycoperdon ericaeum</i>	19	22	·	41	13.7	14	12.5	4	2
<i>Gymnopus androsaceus</i>	20	41	6	67	22.3	21	10.0	5	3
<i>Rhodocybe caelata</i>	4	2	·	6	2.0	2	10.0	3	2
<i>Hygrocybe conica</i> var. <i>conica</i>	·	5	·	5	1.7	5	10.0	1	1
<i>Hygrocybe virginea</i> var. <i>virginea</i>	2	7	1	10	3.3	7	7.5	3	3
<i>Clitocybe vibecina</i>	2	6	·	8	2.7	4	7.5	3	2
<i>Mycena olivaceomarginata</i>	·	8	·	8	2.7	7	7.5	2	1
<i>Entoloma conferendum</i>	·	·	15	15	5.0	15	5.0	1	1
<i>Galerina atkinsoniana</i>	2	·	6	8	2.7	6	5.0	2	2
<i>Clitocybe odora</i> var. <i>odora</i>	·	·	6	6	2.0	6	5.0	1	1
<i>Cystoderma jasonis</i> var. <i>lilacipes</i>	·	5	·	5	1.7	5	5.0	1	1
<i>Flammulaster granulatus</i>	1	1	·	2	0.7	1	5.0	2	2
<i>Mycena galopus</i>	·	10	·	10	3.3	10	2.5	1	1
<i>Lycoperdon nigrescens</i>	·	7	·	7	2.3	7	2.5	1	1
<i>Hygroaster asterosporus</i>	·	6	·	6	2.0	6	2.5	1	1
<i>Cystoderma jasonis</i> var. <i>jasonis</i>	·	·	4	4	1.3	4	2.5	1	1
<i>Gymnopus impudicus</i>	·	4	·	4	1.3	4	2.5	1	1
<i>Lycoperdon molle</i>	4	·	·	4	1.3	4	2.5	1	1
<i>Lycoperdon norvegicum</i>	3	·	1	4	1.3	3	2.5	2	2
<i>Collybia tuberosa</i>	·	2	·	2	0.7	2	2.5	1	1
<i>Mycena stylobates</i>	·	2	·	2	0.7	2	2.5	1	1
<i>Paneolus papilionaceus</i>	·	·	2	2	0.7	2	2.5	1	1
<i>Collybia cookei</i>	1	·	·	1	0.3	1	2.5	1	1
<i>Galerina pumila</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Mycena vitilis</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Polyporus ciliatus</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Tapinella atrotomentosa</i>	·	·	1	1	0.3	1	2.5	1	1

Abbreviations: see Tab. A3.2

Table A3.6  
Macrofungi in plot 7 (phase 6, Jelonka reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Cortinarius cinnamomeoluteus</i>	255	406	277	938	312.7	330	100.0	11	3
<i>Hygrophorus hypothejus</i>	460	357	219	1036	345.3	460	95.0	5	3
<i>Suillus luteus</i>	77	134	105	316	105.3	72	87.5	10	3
<i>Cortinarius</i> spp. I	61	230	99	390	130.0	195	77.5	9	3
<i>Chalciporus piperatus</i>	27	127	58	212	70.7	63	77.5	8	3
<i>Amanita muscaria</i>	72	236	193	501	167.0	235	75.0	6	3
<i>Laccaria proxima</i>	116	298	159	573	191.0	277	62.5	9	3
<i>Tricholoma albobrunneum</i>	3	37	24	64	21.3	37	55.0	4	3
<i>Inocybe lacera</i>	47	51	45	143	47.7	43	50.0	9	3
<i>Lactarius rufus</i>	1	110	51	162	54.0	110	47.5	6	3
<i>Thelephora terrestris</i>	35	38	3	76	25.3	27	37.5	6	3
<i>Cortinarius colus</i>	·	56	·	56	18.7	56	35.0	1	1
<i>Tricholoma equestre</i>	3	38	34	75	25.0	37	32.5	6	3
<i>Rhodocollybia butyracea</i> f. <i>asema</i>	2	6	31	39	13.0	26	32.5	6	3
<i>Tricholoma portentosum</i>	8	21	49	78	26.0	45	30.0	4	3
<i>Laccaria bicolor</i>	58	25	93	176	58.7	58	25.0	8	3
<i>Leccinum aurantiacum</i>	1	8	12	21	7.0	9	17.5	6	3
<i>Russula fragilis</i>	2	48	4	54	18.0	45	20.0	5	3

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Cortinarius parevernus</i>	·	10	11	21	7.0	11	15.0	2	2
<i>Cortinarius</i> spp. III	2	9	6	17	5.7	9	15.0	3	3
<i>Boletus edulis</i>	·	·	7	7	2.3	3	15.0	3	1
<i>Inocybe curvipes</i>	·	19	·	19	6.3	19	10.0	1	1
<i>Tricholoma saponaceum</i> var. <i>saponaceum</i>	·	2	15	17	5.7	15	10.0	2	2
<i>Cortinarius</i> spp. II		1	12	13	4.3	8	10.0	3	2
<i>Ramaria eumorpha</i>	19	31	9	59	19.7	31	7.5	3	3
<i>Amanita rubescens</i>	·	3	2	5	1.7	3	7.5	2	2
<i>Cortinarius mucosus</i>	·	3	·	3	1.0	3	7.5	1	1
<i>Tricholoma pessundatum</i>	3	·	·	3	1.0	3	7.5	1	1
<i>Cortinarius purpurascens</i>	·	·	4	4	1.3	4	5.0	1	1
<i>Inocybe assimilata</i>	·	·	3	3	1.0	3	5.0	1	1
<i>Amanita pantherina</i>	1	1	·	2	0.7	1	5.0	2	2
<i>Tricholoma imbricatum</i>	·	·	3	3	1.0	3	2.5	1	1
<i>Xerocomus badius</i>	·	·	3	3	1.0	3	2.5	1	1
<i>Inocybe calida</i>	2	·	·	2	0.7	2	2.5	1	1
<i>Rhizopogon luteolus</i>	2	·	·	2	0.7	2	2.5	1	1
<i>Clitopilus prunulus</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Inocybe asterospora</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Paxillus involutus</i>	·	·	1	1	0.3	1	2.5	1	1
<b>Other species</b>									
<i>Clitocybe diatreta</i>	345	239	141	725	241.7	208	85.0	8	3
<i>Psilocybe montana</i>	500	295	54	849	283.0	254	80.0	14	3
<i>Rickenella fibula</i>	45	156	82	283	94.3	98	65.0	10	3
<i>Cystoderma amianthinum</i>	62	130	173	365	121.7	131	62.5	10	3
<i>Galerina atkinsoniana</i>	50	185	132	367	122.3	159	50.0	6	3
<i>Gymnopus androsaceus</i>	252	14	87	353	117.7	247	47.5	7	3
<i>Ampulloclitocybe clavipes</i>	29	54	25	108	36.0	45	45.0	6	3
<i>Galerina</i> spp.	130	64	39	233	77.7	113	40.0	6	3
<i>Hygroaster asterosporus</i>	10	17	34	61	20.3	23	40.0	4	3
<i>Mycena leptcephala</i>	95	59	7	161	53.7	92	37.5	6	3
<i>Collybia cirrhata</i>	64	20	23	107	35.7	40	32.5	4	3
<i>Entoloma</i> spp.	5	13	18	36	12.0	13	30.0	6	3
<i>Cantharellula umbonata</i>	74	122	138	334	111.3	65	22.5	10	3
<i>Clitocybe vibecina</i>	16	41	4	61	20.3	31	22.5	5	3
<i>Mycena pura</i>	19	2	1	22	7.3	19	20.0	4	3
<i>Clitocybe candicans</i>	22	2	·	24	8.0	20	17.5	4	2
<i>Lycoperdon perlatum</i>	6	4	5	15	5.0	4	15.0	5	3
<i>Strobilurus stephanocystis</i>	·	10	4	14	4.7	10	15.0	2	2
<i>Flammulaster granulatus</i>	9	25	·	34	11.3	25	12.5	2	2
<i>Cystoderma granulosa</i>	18	3	4	25	8.3	17	12.5	5	3
<i>Cystoderma jasonis</i> var. <i>lilacipes</i>	9	·	10	19	6.3	9	12.5	3	2
<i>Dacrymyces variisporus</i>	·	60	·	60	20.0	37	10.0	3	1
<i>Mycena vitilis</i>	·	4	5	9	3.0	5	10.0	2	2
<i>Rhodocybe caelata</i>	4	·	1	5	1.7	3	10.0	3	2
<i>Clavaria argillacea</i>	2	15	·	17	5.7	15	7.5	2	2
<i>Mycena flavoalba</i>	4	11	·	15	5.0	10	7.5	3	2
<i>Mycena stylobates</i>	2	1	1	4	1.3	2	7.5	3	3
<i>Marasmiellus vaillantii</i>	·	25	30	55	18.3	30	5.0	2	2
<i>Galerina hypnorum</i>	9	20	·	29	9.7	16	5.0	3	2
<i>Crinipellis scabella</i>	·	·	18	18	6.0	18	5.0	1	1
<i>Galerina pumila</i>	3	6	·	9	3.0	4	5.0	4	2

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Hygrocybe conica</i> var. <i>conica</i>	·	6	·	6	2.0	6	5.0	1	1
<i>Rhodocybe parilis</i>	2	4	·	6	2.0	4	5.0	2	2
<i>Entoloma jubatum</i>	·	5	·	5	1.7	5	5.0	1	1
<i>Baeospora myosura</i>	·	4	·	4	1.3	4	5.0	1	1
<i>Ripartites tricholoma</i>	1	2	·	3	1.0	2	5.0	2	2
<i>Melanoleuca graminicola</i>	·	2	·	2	0.7	2	5.0	1	1
<i>Mycena vulgaris</i>	25	·	50	75	25.0	50	2.5	2	2
<i>Trichaptum fuscoviolaceum</i>	39	·	·	39	13.0	17	2.5	3	1
<i>Clitocybe fragrans</i>	28	·	·	28	9.3	28	2.5	1	1
<i>Gymnopus aquosus</i>	·	·	11	11	3.7	6	2.5	2	1
<i>Tremella mesenterica</i>	·	·	9	9	3.0	4	2.5	3	1
<i>Pholiota gummosa</i>	·	4	4	8	2.7	4	2.5	2	2
<i>Calocera furcata</i>	5	·	·	5	1.7	5	2.5	1	1
<i>Mycena metata</i>	5	·	·	5	1.7	5	2.5	1	1
<i>Sphaerobolus stellatus</i>	·	5	·	5	1.7	5	2.5	1	1
<i>Collybia tuberosa</i>	·	3	·	3	1.0	3	2.5	1	1
<i>Phanerochaete sanguinea</i>	3	·	·	3	1.0	3	2.5	1	1
<i>Leptosporomyces galzinii</i>	·	2	·	2	0.7	2	2.5	1	1
<i>Mycena sanguinolenta</i>	2	·	·	2	0.7	2	2.5	1	1
<i>Galerina badipes</i>	1	·	·	1	0.3	1	2.5	1	1
<i>Gymnopus putillus</i>	1	·	·	1	0.3	1	2.5	1	1
<i>Hypholoma fasciculare</i>	·	1	·	1	0.3	1	2.5	1	1

Abbreviations: see Tab. A3.2

Table A3.7  
Macrofungi in plot 13 (phase 6, Jelonka reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Cortinarius</i> spp. I	595	1204	445	2244	748.0	859	100.0	10	3
<i>Hygrophorus hypothejus</i>	437	443	265	1145	381.7	437	97.5	6	3
<i>Cortinarius cinnamomeoluteus</i>	185	427	180	792	264.0	349	95.0	11	3
<i>Amanita muscaria</i>	69	194	162	425	141.7	186	90.0	8	3
<i>Suillus luteus</i>	26	155	61	242	80.7	79	90.0	8	3
<i>Tricholoma albobrunneum</i>	34	132	72	238	79.3	105	75.0	5	3
<i>Laccaria proxima</i>	89	325	166	580	193.3	276	72.5	6	3
<i>Chalciporus piperatus</i>	7	105	58	170	56.7	65	72.5	6	3
<i>Tricholoma portentosum</i>	35	138	10	183	61.0	136	52.5	5	3
<i>Cortinarius parevemi</i>	1	51	88	140	46.7	70	52.5	4	3
<i>Thelephora terrestris</i>	96	123	57	276	92.0	109	50.0	11	3
<i>Cortinarius mucosus</i>	7	18	26	51	17.0	25	47.5	6	3
<i>Inocybe lacera</i>	34	51	31	116	38.7	45	45.0	8	3
<i>Rhodocollybia butyracea</i> f. <i>asema</i>	18	17	30	65	21.7	30	37.5	4	3
<i>Tricholoma equestre</i>	11	25	54	90	30.0	47	35.0	5	3
<i>Cortinarius anthracinus</i>	3	26	1	30	10.0	26	32.5	4	3
<i>Tricholoma saponaceum</i> var. <i>saponaceum</i>	29	45	50	124	41.3	45	25.0	4	3
<i>Cortinarius multififormis</i> f. <i>coniferarum</i>	2	5	14	21	7.0	10	17.5	4	3
<i>Lactarius rufus</i>	·	12	1	13	4.3	12	15.0	2	2
<i>Sarcodon squamosus</i>	5	8	·	13	4.3	8	15.0	3	2
<i>Laccaria bicolor</i>	22	9	20	51	17.0	22	12.5	6	3
<i>Hebeloma</i> sp. 1	3	9	·	12	4.0	9	10.0	2	2
<i>Ramaria eumorpha</i>	7	21	·	28	9.3	21	7.5	2	2

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Cortinarius</i> spp. III	·	5	·	5	1.7	5	7.5	1	1
<i>Clitopilus prunulus</i>	17	15	·	32	10.7	17	5.0	2	2
<i>Cortinarius</i> spp. II	7	1	·	8	2.7	7	5.0	2	2
<i>Inocybe assimilata</i>	1	3	·	4	1.3	2	5.0	3	2
<i>Cortinarius purpurascens</i>	·	3	·	3	1.0	3	5.0	1	1
<i>Inocybe praetervisa</i>	·	2	1	3	1.0	2	5.0	2	2
<i>Amanita pantherina</i>	·	1	1	2	0.7	1	5.0	2	2
<i>Suillus bovinus</i>	·	1	1	2	0.7	1	2.5	2	2
<i>Boletus edulis</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Cortinarius acetosus</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Cortinarius cinnabarinus</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Cortinarius</i> spp. IV	·	1	·	1	0.3	1	2.5	1	1
<i>Inocybe curvipes</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Paxillus involutus</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Tricholoma imbricatum</i>	·	1	·	1	0.3	1	2.5	1	1
<b>Other species</b>									
<i>Psilocybe montana</i>	177	91	25	293	97.7	92	95.0	11	3
<i>Clitocybe diatreta</i>	606	606	206	1418	472.7	341	90.0	11	3
<i>Cystoderma amianthinum</i>	78	129	31	238	79.3	106	65.0	9	3
<i>Mycena leptcephala</i>	44	63	11	118	39.3	59	60.0	5	3
<i>Rickenella fibula</i>	23	337	19	379	126.3	175	52.5	8	3
<i>Galerina atkinsoniana</i>	20	222	60	302	100.7	162	52.5	7	3
<i>Galerina</i> spp.	32	144	19	195	65.0	85	50.0	9	3
<i>Gymnopus androsaceus</i>	328	34	5	367	122.3	238	37.5	5	3
<i>Entoloma</i> spp.	7	16	6	29	9.7	12	35.0	6	3
<i>Clavaria argillacea</i>	67	177	23	267	89.0	160	32.5	5	3
<i>Hygroaster asterosporus</i>	5	15	2	22	7.3	15	30.0	3	3
<i>Collybia cirrhata</i>	63	56	37	156	52.0	56	27.5	5	3
<i>Galerina pumila</i>	5	14	·	19	6.3	7	15.0	3	2
<i>Lycoperdon perlatum</i>	1	28	·	29	9.7	22	12.5	3	2
<i>Ampulloclitocybe clavipes</i>	2	10	2	14	4.7	9	10.0	5	3
<i>Melanoleuca graminicola</i>	4	9	·	13	4.3	5	10.0	4	2
<i>Mycena flavoalba</i>	3	10	·	13	4.3	9	10.0	3	2
<i>Clitocybe candicans</i>	10	2	7	19	6.3	10	7.5	4	3
<i>Lycoperdon molle</i>	6	3	5	14	4.7	6	7.5	5	3
<i>Dacrymyces variisporus</i>	5	3	3	11	3.7	5	7.5	3	3
<i>Marasmius oreades</i>	6	3	1	10	3.3	3	7.5	4	3
<i>Entoloma jubatum</i>	·	4	3	7	2.3	4	7.5	2	2
<i>Mycena vitilis</i>	5	2	·	7	2.3	4	7.5	3	2
<i>Lycoperdon excipuliforme</i>	4	1	·	5	1.7	2	7.5	3	2
<i>Collybia tuberosa</i>	40	·	5	45	15.0	32	5.0	3	2
<i>Cystoderma jasonis</i> var. <i>lilacipes</i>	·	·	9	9	3.0	9	5.0	1	1
<i>Mycena olivaceomarginata</i>	8	·	·	8	2.7	8	5.0	1	1
<i>Leptosporomyces galzinii</i>	·	5	·	5	1.7	5	5.0	1	1
<i>Rhodocybe caelata</i>	1	1	2	4	1.3	2	5.0	3	3
<i>Rhodocybe parilis</i>	1	2	·	3	1.0	2	5.0	2	2
<i>Mycena parabolica</i>	2	·	·	2	0.7	2	5.0	1	1
<i>Ripartites tricholoma</i>	2	·	·	2	0.7	2	5.0	1	1
<i>Gymnopus confluens</i>	·	3	14	17	5.7	14	2.5	2	2
<i>Rhodocybe hirneola</i>	·	12	·	12	4.0	12	2.5	1	1
<i>Cystodermella granulosa</i>	4	·	·	4	1.3	4	2.5	1	1
<i>Clitocybe vibecina</i>	·	3	·	3	1.0	3	2.5	1	1
<i>Gymnopus putillus</i>	3	·	·	3	1.0	3	2.5	1	1

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Lycoperdon dermoxanthum</i>	1	2	·	3	1.0	2	2.5	2	2
<i>Mycena speirea</i>	·	·	3	3	1.0	3	2.5	1	1
<i>Lepiota pseudolilacea</i>	·	2	·	2	0.7	2	2.5	1	1
<i>Entoloma nitens</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Hygrocybe virginea</i> var. <i>virginea</i>	1	·	·	1	0.3	1	2.5	1	1
<i>Mycena arcangeliana</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Psathyrella fibrillosa</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Strobilurus stephanocystis</i>	·	·	1	1	0.3	1	2.5	1	1

Abbreviations: see Tab. A3.2

Table A3.8  
Macrofungi in plot 1 (phase 6/7, Jelonka reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Cortinarius</i> spp. I	286	464	218	968	322.7	304	95.0	8	3
<i>Amanita muscaria</i>	70	150	101	321	107.0	148	90.0	5	3
<i>Cortinarius cinnamomeoluteus</i>	384	470	162	1016	338.7	345	87.5	10	3
<i>Hygrophorus hypothejus</i>	313	131	95	539	179.7	313	77.5	5	3
<i>Tricholoma albobrunneum</i>	28	130	18	176	58.7	97	75.0	5	3
<i>Tricholoma equestre</i>	62	189	56	307	102.3	168	72.5	8	3
<i>Chalciporus piperatus</i>	3	71	15	89	29.7	51	72.5	7	3
<i>Laccaria proxima</i>	101	194	91	386	128.7	144	57.5	9	3
<i>Suillus luteus</i>	22	49	61	132	44.0	38	57.5	8	3
<i>Lactarius rufus</i>	9	343	110	462	154.0	331	52.5	8	3
<i>Tricholoma portentosum</i>	81	120	21	222	74.0	115	52.5	5	3
<i>Rhodocollybia butyracea</i> f. <i>asema</i>	14	14	62	90	30.0	47	40.0	5	3
<i>Thelephora terrestris</i>	27	68	5	100	33.3	53	35.0	9	3
<i>Cortinarius parevernus</i>	·	18	38	56	18.7	25	35.0	3	2
<i>Tricholoma saponaceum</i> var. <i>saponaceum</i>	30	51	58	139	46.3	51	30.0	4	3
<i>Inocybe lacera</i>	22	84	24	130	43.3	63	30.0	9	3
<i>Boletus edulis</i>	·	9	7	16	5.3	5	27.5	5	2
<i>Cortinarius</i> spp. II	6	17	·	23	7.7	17	20.0	2	2
<i>Cortinarius mucosus</i>	1	3	12	16	5.3	12	15.0	3	3
<i>Cortinarius</i> spp. III	2	6	·	8	2.7	6	10.0	2	2
<i>Cortinarius anthracinus</i>	·	5	2	7	2.3	5	10.0	2	2
<i>Cortinarius purpurascens</i>	1	1	2	4	1.3	1	10.0	4	3
<i>Sarcodon squamosus</i>	5	7	2	14	4.7	7	7.5	3	3
<i>Inocybe assimolata</i>	·	4	1	5	1.7	4	7.5	2	2
<i>Xerocomus badius</i>	·	4	1	5	1.7	3	7.5	3	2
<i>Clitopilus prunulus</i>	1	2	1	4	1.3	2	5.0	3	3
<i>Amanita rubescens</i>	·	1	1	2	0.7	1	5.0	2	2
<i>Boletus reticulatus</i>	·	2	·	2	0.7	2	5.0	1	1
<i>Laccaria bicolor</i>	·	·	15	15	5.0	15	2.5	1	1
<i>Cortinarius sanguineus</i>	·	6	·	6	2.0	6	2.5	1	1
<i>Cortinarius multififormis</i> f. <i>coniferarum</i>	·	1	2	3	1.0	2	2.5	2	2
<i>Paxillus involutus</i>	·	2	·	2	0.7	2	2.5	1	1
<i>Boletus pinophilus</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Leccinum aurantiacum</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Suillus bovinus</i>	·	1	·	1	0.3	1	2.5	1	1



Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>Other species</b>									
<i>Clitocybe diatreta</i>	447	561	202	1210	403.3	279	97.5	9	3
<i>Psilocybe montana</i>	197	114	31	342	114.0	111	72.5	11	3
<i>Gymnopus androsaceus</i>	694	227	142	1063	354.3	564	67.5	9	3
<i>Mycena leptocephala</i>	88	25	10	123	41.0	71	57.5	5	3
<i>Cystoderma amianthinum</i>	71	54	10	135	45.0	65	52.5	7	3
<i>Galerina</i> spp.	126	161	9	296	98.7	117	50.0	6	3
<i>Galerina atkinsoniana</i>	30	263	42	335	111.7	186	42.5	6	3
<i>Collybia cirrhata</i>	14	45	69	128	42.7	49	27.5	7	3
<i>Mycena speirea</i>	90	.	.	90	30.0	90	27.5	1	1
<i>Hygroaster asterosporus</i>	7	21	10	38	12.7	21	27.5	4	3
<i>Entoloma</i> spp.	12	17	8	37	12.3	9	27.5	6	3
<i>Mycena vitilis</i>	15	11	3	29	9.7	9	27.5	6	3
<i>Galerina hypnorum</i>	16	50	7	73	24.3	47	25.0	5	3
<i>Galerina pumila</i>	1	42	8	51	17.0	33	25.0	6	3
<i>Clavaria argillacea</i>	105	266	63	434	144.7	259	22.5	5	3
<i>Mycena flavoalba</i>	19	12	1	32	10.7	18	20.0	5	3
<i>Rickenella fibula</i>	11	24	31	66	22.0	16	17.5	5	3
<i>Dacrymyces variisporus</i>	.	96	.	96	32.0	96	15.0	1	1
<i>Strobilurus stephanocystis</i>	3	51	19	73	24.3	51	15.0	3	3
<i>Clitocybe vibecina</i>	3	19	.	22	7.3	19	15.0	2	2
<i>Clitocybe phyllophila</i>	1	41	8	50	16.7	41	12.5	3	3
<i>Ampulloclitocybe clavipes</i>	.	9	3	12	4.0	8	12.5	5	2
<i>Mycena olivaceomarginata</i>	8	.	1	9	3.0	7	12.5	3	2
<i>Ripartites tricholoma</i>	.	7	.	7	2.3	7	12.5	1	1
<i>Cystoderma jasonis</i> var. <i>jasonis</i>	9	13	.	22	7.3	13	10.0	2	2
<i>Clitocybe anisata</i>	6	8	.	14	4.7	8	10.0	2	2
<i>Collybia tuberosa</i>	2	4	1	7	2.3	3	10.0	4	3
<i>Leptosporomyces galzinii</i>	.	11	.	11	3.7	11	7.5	1	1
<i>Cystoderma jasonis</i> var. <i>lilacipes</i>	5	5	.	10	3.3	5	7.5	2	2
<i>Clitocybe fragrans</i>	1	7	.	8	2.7	7	7.5	2	2
<i>Mycena stylobates</i>	6	.	.	6	2.0	6	7.5	1	1
<i>Dacrymyces stillatus</i>	10	20	.	30	10.0	20	5.0	2	2
<i>Lycoperdon perlatum</i>	14	8	.	22	7.3	14	5.0	4	2
<i>Galerina mniophila</i>	15	.	.	15	5.0	15	5.0	1	1
<i>Gymnopus impudicus</i>	13	.	.	13	4.3	13	5.0	1	1
<i>Marasmius oreades</i>	7	.	.	7	2.3	4	5.0	2	1
<i>Amylostereum laevigatum</i>	.	5	.	5	1.7	5	5.0	1	1
<i>Auriscalpium vulgare</i>	1	3	.	4	1.3	3	5.0	2	2
<i>Hygrocybe conica</i> var. <i>conica</i>	.	2	.	2	0.7	2	5.0	1	1
<i>Panellus mitis</i>	.	6	.	6	2.0	6	2.5	1	1
<i>Cystodermella granulosa</i>	.	.	5	5	1.7	5	2.5	1	1
<i>Hygrocybe virginea</i> var. <i>virginea</i>	.	3	.	3	1.0	3	2.5	1	1
<i>Gymnopus putillus</i>	2	.	.	2	0.7	2	2.5	1	1
<i>Mycena pura</i>	.	2	.	2	0.7	2	2.5	1	1
<i>Agaricus dulcidulus</i>	1	.	.	1	0.3	1	2.5	1	1
<i>Baeospora myosura</i>	.	.	1	1	0.3	1	2.5	1	1
<i>Cystoderma carcharias</i>	1	.	.	1	0.3	1	2.5	1	1
<i>Entoloma undatum</i>	.	1	.	1	0.3	1	2.5	1	1
<i>Hypholoma lateritium</i>	.	.	1	1	0.3	1	2.5	1	1
<i>Mycena aetites</i>	1	.	.	1	0.3	1	2.5	1	1

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Mycena septentrionalis</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Phanerochaete sanguinea</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Psathyrella fibrillosa</i>	·	1	·	1	0.3	1	2.5	1	1

Abbreviations: see Tab. A3.2

Table A3.9  
Macrofungi in plot 6 (phase 6/7, Jelonka reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Cortinarius cinnamomeoluteus</i>	211	431	271	913	304.3	336	97.5	11	3
<i>Cortinarius</i> spp. I	100	382	246	728	242.7	271	97.5	8	3
<i>Amanita muscaria</i>	99	153	155	407	135.7	152	95.0	7	3
<i>Hygrophorus hypothejus</i>	337	166	76	579	193.0	337	92.5	5	3
<i>Tricholoma albobrunneum</i>	25	177	62	264	88.0	167	90.0	5	3
<i>Tricholoma portentosum</i>	71	141	73	285	95.0	141	77.5	4	3
<i>Laccaria proxima</i>	173	286	183	642	214.0	257	72.5	7	3
<i>Suillus luteus</i>	17	109	69	195	65.0	79	72.5	8	3
<i>Tricholoma equestre</i>	30	113	47	190	63.3	107	72.5	8	3
<i>Chalciporus piperatus</i>	4	50	21	75	25.0	27	67.5	7	3
<i>Lactarius rufus</i>	2	131	33	166	55.3	125	42.5	8	3
<i>Tricholoma saponaceum</i> var. <i>saponaceum</i>	17	44	67	128	42.7	48	37.5	5	3
<i>Cortinarius anthracinus</i>	·	38	·	38	12.7	38	37.5	1	1
<i>Cortinarius parevemi</i>	·	46	18	64	21.3	46	35.0	3	2
<i>Laccaria bicolor</i>	·	47	35	82	27.3	20	32.5	6	2
<i>Rhodocollybia butyracea</i> f. <i>asema</i>	8	10	22	40	13.3	16	32.5	6	3
<i>Boletus edulis</i>	7	4	14	25	8.3	7	32.5	7	3
<i>Cortinarius</i> spp. II	2	16	30	48	16.0	22	20.0	4	3
<i>Sarcodon squamosus</i>	3	9	4	16	5.3	9	15.0	3	3
<i>Cortinarius mucosus</i>	·	1	7	8	2.7	7	15.0	2	2
<i>Thelephora terrestris</i>	19	9	3	31	10.3	19	12.5	3	3
<i>Inocybe lacera</i>	·	14	·	14	4.7	11	12.5	2	1
<i>Cortinarius purpurascens</i>	·	7	·	7	2.3	7	12.5	1	1
<i>Clitopilus prunulus</i>	·	6	·	6	2.0	5	10.0	2	1
<i>Inocybe assimilata</i>	3	20	·	23	7.7	20	7.5	2	2
<i>Cortinarius</i> spp. IV	·	·	17	17	5.7	17	5.0	1	1
<i>Amanita rubescens</i>	·	3	3	6	2.0	3	5.0	3	2
<i>Amanita pantherina</i>	·	4	·	4	1.3	4	5.0	1	1
<i>Inocybe curvipes</i>	·	3	1	4	1.3	3	5.0	2	2
<i>Suillus bovinus</i>	·	4	·	4	1.3	4	5.0	1	1
<i>Rhizopogon luteolus</i>	·	1	1	2	0.7	1	5.0	2	2
<i>Ramaria eumorpha</i>	5	·	·	5	1.7	5	2.5	1	1
<i>Cortinarius</i> spp. III	2	·	·	2	0.7	2	2.5	1	1
<i>Boletus pinophilus</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Cortinarius multififormis</i> f. <i>coniferarum</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Inocybe calida</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Inocybe praetervisa</i>	·	1	·	1	0.3	1	2.5	1	1
<b>Other species</b>									
<i>Clitocybe diatreta</i>	280	87	42	409	136.3	193	97.5	9	3
<i>Psilocybe montana</i>	207	229	35	471	157.0	136	95.0	12	3

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Clitocybe vibecina</i>	17	103	132	252	84	130	30.0	4	3
<i>Strobilurus stephanocystis</i>	4	19	26	49	16.3	23	30.0	4	3
<i>Mycena leptcephala</i>	3	39	·	42	14.0	39	25.0	2	2
<i>Gymnopus androsaceus</i>	119	19	26	164	54.7	119	22.5	4	3
<i>Collybia cirrhata</i>	14	30	102	146	48.7	52	22.5	5	3
<i>Galerina atkinsoniana</i>	6	40	·	46	15.3	38	22.5	4	2
<i>Hygroaster asterosporus</i>	30	3	·	33	11.0	30	22.5	2	2
<i>Clavaria argillacea</i>	102	21	·	123	41.0	102	20.0	2	2
<i>Leptosporomyces galzinii</i>	·	37	·	37	12.3	37	20.0	1	1
<i>Cystoderma amianthinum</i>	3	17	3	23	7.7	10	17.5	6	3
<i>Ampulloclitocybe clavipes</i>	6	14	·	20	6.7	13	17.5	4	2
<i>Galerina</i> spp.	10	51	1	62	20.7	49	15.0	4	3
<i>Macrolepiota procera</i>	3	4	·	7	2.3	4	15.0	3	2
<i>Entoloma</i> spp.	12	1	·	13	4.3	12	12.5	2	2
<i>Baeospora myosura</i>	2	5	1	8	2.7	3	12.5	4	3
<i>Mycena vitilis</i>	3	1	·	4	1.3	3	7.5	2	2
<i>Lycoperdon perlatum</i>	·	7	6	13	4.3	7	5.0	2	2
<i>Dacrymyces variisporus</i>	2	7	·	9	3.0	7	5.0	2	2
<i>Clitocybe fragrans</i>	·	·	7	7	2.3	7	5.0	1	1
<i>Rickenella fibula</i>	·	7	·	7	2.3	7	5.0	1	1
<i>Mycena pura</i>	6	·	·	6	2.0	3	5.0	2	1
<i>Rhodocybe caelata</i>	1	1	·	2	0.7	1	5.0	2	2
<i>Hygrocybe conica</i> var. <i>conica</i>	·	3	·	3	1.0	3	2.5	1	1
<i>Hypholoma fasciculare</i>	·	·	3	3	1.0	3	2.5	1	1
<i>Hygrophoropsis aurantiaca</i>	·	·	2	2	0.7	2	2.5	1	1
<i>Hypholoma lateritium</i>	·	2	·	2	0.7	2	2.5	1	1
<i>Mycena metata</i>	2	·	·	2	0.7	2	2.5	1	1
<i>Auriscalpium vulgare</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Clitocybe candicans</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Cystoderma carcharias</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Flammulaster granulosus</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Gymnopus putillus</i>	1	·	·	1	0.3	1	2.5	1	1
<i>Mycena aetites</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Mycena flavoalba</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Mycena stylobates</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Phanerochaete sanguinea</i>	·	1	·	1	0.3	1	2.5	1	1
<i>Polyporus ciliatus</i>	1	·	·	1	0.3	1	2.5	1	1
<i>Rhodocybe parilis</i>	·	·	1	1	0.3	1	2.5	1	1
<i>Tapinella atrotomentosa</i>	·	·	1	1	0.3	1	2.5	1	1

Abbreviations: see Tab. A3.2

Table A3.10  
Macrofungi in plot 11 (phase 8, Jelonka reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Rhodocollybia butyracea</i> f. <i>asema</i>	54	14	78	146	48.7	55	100.0	6	3
<i>Cortinarius</i> spp. II	42	416	31	489	163.0	416	93.3	4	3
<i>Cortinarius trivialis</i>	18	169	27	214	71.3	167	86.7	7	3
<i>Cortinarius</i> spp. I	5	97	86	188	62.7	84	86.7	8	3
<i>Amanita citrina</i>	1	98	2	101	33.7	95	86.7	5	3
<i>Hebeloma</i> sp. 3	4	29	65	98	32.7	65	73.3	4	3
<i>Leccinum aurantiacum</i>	4	14	30	48	16.0	29	73.3	7	3
<i>Russula clavipes</i>	7	17	20	44	14.7	17	73.3	5	3

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Cortinarius</i> spp. III	1	46	37	84	28.0	46	66.7	3	3
<i>Amanita muscaria</i>	25	19	29	73	24.3	28	60.0	5	3
<i>Tricholoma equestre</i>	4	14	1	19	6.3	11	53.3	4	3
<i>Cortinarius alboviolaceus</i>		73	10	83	27.7	53	40.0	4	2
<i>Russula fragilis</i>	2	8	2	12	4.0	8	40.0	4	3
<i>Suillus luteus</i>	·	6	11	17	5.7	11	33.3	2	2
<i>Clitopilus prunulus</i>	·	6	1	7	2.3	4	33.3	3	2
<i>Amanita gemmata</i>	3	2	·	5	1.7	3	33.3	2	2
<i>Paxillus involutus</i>	1	19	15	35	11.7	19	26.7	6	3
<i>Tricholoma portentosum</i>	5	5	3	13	4.3	5	26.7	3	3
<i>Boletus edulis</i>	1	3	3	7	2.3	2	26.7	5	3
<i>Chalciporus piperatus</i>	·	2	3	5	1.7	2	26.7	3	2
<i>Russula galochroa</i>	·	2	3	5	1.7	2	26.7	3	2
<i>Thelephora caryophylla</i>	3	3	·	6	2.0	3	20.0	2	2
<i>Hygrophorus hypothejus</i>	4	1	·	5	1.7	4	20.0	2	2
<i>Amanita pantherina</i>	1	3	·	4	1.3	3	20.0	2	2
<i>Tricholoma imbricatum</i>	·	19	·	19	6.3	19	13.3	1	1
<i>Lactarius uvidus</i>	·	10	·	10	3.3	10	13.3	1	1
<i>Cortinarius cinnamomeoluteus</i>	3	·	·	3	1.0	3	13.3	1	1
<i>Lactarius rufus</i>	2	1	·	3	1.0	2	13.3	2	2
<i>Cortinarius parevernus</i>	·	2	·	2	0.7	2	13.3	1	1
<i>Lactarius necator</i>	·	·	3	3	1.0	3	6.7	1	1
<i>Lactarius torminosus</i>	1	2	·	3	1.0	2	6.7	2	2
<i>Amanita rubescens</i>	·	1	1	2	0.7	1	6.7	2	2
<i>Laccaria proxima</i>	·	·	1	1	0.3	1	6.7	1	1
<i>Thelephora terrestris</i>	·	1	·	1	0.3	1	6.7	1	1
<b>Other species</b>									
<i>Auriscalpium vulgare</i>	41	12	13	66	22.0	38	86.7	7	3
<i>Lycoperdon molle</i>	22	15	25	62	20.7	16	80.0	10	3
<i>Clitocybe candicans</i>	49	4	·	53	17.7	28	80.0	3	2
<i>Psathyrella candolleana</i>	59	24	2	85	28.3	57	73.3	6	3
<i>Galerina hypnorum</i>	19	63	15	97	32.3	55	66.7	5	3
<i>Mycena leptcephala</i>	67	17	6	90	30.0	66	66.7	7	3
<i>Gymnopus androsaceus</i>	183	223	51	457	152.3	214	60.0	10	3
<i>Tubaria furfuracea</i>	83	128	14	225	75.0	109	60.0	7	3
<i>Mycena pura</i>	52	6	1	59	19.7	52	60.0	4	3
<i>Lycoperdon perlatum</i>	7	14	3	24	8.0	13	60.0	5	3
<i>Inonotus rheades</i>	28	·	9	x	x	28	60.0	8	2
<i>Dacrymyces stillatus</i>	100	1090	135	1325	441.7	1080	53.3	5	3
<i>Cystoderma carcharias</i>	13	29	29	71	23.7	29	53.3	4	3
<i>Clitocybe phyllophila</i>	15	48	1	64	21.3	45	53.3	4	3
<i>Entoloma</i> spp.	7	37	·	44	14.7	30	53.3	3	2
<i>Baeospora myosura</i>	9	27	·	36	12.0	27	53.3	3	2
<i>Dacrymyces variisporus</i>	10	142	40	192	64.0	137	46.7	5	3
<i>Ampulloclitocybe clavipes</i>	11	21	4	36	12.0	21	46.7	4	3
<i>Gymnopus confluens</i>	301	231	280	812	270.7	255	40.0	9	3
<i>Mycena sanguinolenta</i>	2	44	·	46	15.3	44	40.0	2	2
<i>Clitocybe diatreta</i>	39	3	3	45	15.0	22	40.0	4	3
<i>Galerina marginata</i>	4	34	·	38	12.7	34	40.0	3	2
<i>Flammulaster granulosus</i>	28	2	2	32	10.7	28	40.0	4	3
<i>Crucibulum laeve</i>	1	10	17	28	9.3	12	40.0	4	3
<i>Lycoperdon nigrescens</i>	5	11	3	19	6.3	11	40.0	5	3
<i>Mycena olivaceomarginata</i>	8	4	·	12	4.0	8	40.0	3	2

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Peniophora polygonia</i>	110	94	35	239	79.7	110	33.3	5	3
<i>Flammulaster rhombisporus</i>	2	57	18	77	25.7	49	33.3	4	3
<i>Mycena stipata</i>	22	·	·	22	7.3	20	33.3	2	1
<i>Clitocybe fragrans</i>	·	7	3	10	3.3	6	33.3	4	2
<i>Collybia cirrhata</i>	16	154	30	200	66.7	154	26.7	4	3
<i>Stereum hirsutum</i>	38	25	30	93	31.0	28	26.7	6	3
<i>Xeromphalina cornui</i>	42	25	·	67	22.3	35	26.7	3	2
<i>Amylostereum laevigatum</i>	·	56	·	56	18.7	56	26.7	1	1
<i>Gymnopus aquosus</i>	13	13	27	53	17.7	27	26.7	3	3
<i>Clitocybe metachroa</i>	20	19	3	42	14.0	20	26.7	3	3
<i>Mycena galopus</i>	1	12	·	13	4.3	11	26.7	3	2
<i>Mycena stylobates</i>	4	3	2	9	3.0	4	26.7	3	3
<i>Mycena rubromarginata</i>	5	·	·	5	1.7	5	26.7	1	1
<i>Macrolepiota procera</i>	4	·	·	4	1.3	4	26.7	1	1
<i>Daedaleopsis confragosa</i>	4	2	6	x	x	6	26.7	7	3
<i>Sphaerobolus stellatus</i>	109	12	·	121	40.3	60	20.0	3	2
<i>Gleocystidiellum porosum</i>	50	2	·	52	17.3	30	20.0	3	2
<i>Mycena filopes</i>	24	·	3	27	9.0	23	20.0	3	2
<i>Exidia plana</i>	3	10	10	23	7.7	10	20.0	3	3
<i>Clitocybe odora</i> var. <i>odora</i>	17	2	·	19	6.3	17	20.0	2	2
<i>Strobilurus tenacellus</i>	7	1	·	8	2.7	7	20.0	2	2
<i>Mycena vitilis</i>	6	·	·	6	2.0	6	20.0	1	1
<i>Conocybe ochrostriata</i>	·	3	·	3	1.0	3	20.0	1	1
<i>Crepidotus lundellii</i>	285	7	20	312	104.0	255	13.3	4	3
<i>Marasmius epiphyllus</i>	·	15	35	50	16.7	35	13.3	2	2
<i>Peniophorella praetermissa</i>	25	4	·	29	9.7	25	13.3	2	2
<i>Phlebia rufa</i>	18	11	·	29	9.7	14	13.3	4	2
<i>Marasmiellus vaillantii</i>	2	·	20	22	7.3	20	13.3	2	2
<i>Phaeomarasmium erinaceus</i>	·	18	2	20	6.7	18	13.3	2	2
<i>Vesiculomyces citrinus</i>	7	13	·	20	6.7	13	13.3	2	2
<i>Tremella mesenterica</i>	·	17	·	17	5.7	17	13.3	1	1
<i>Marasmius oreades</i>	12	·	·	12	4.0	9	13.3	2	1
<i>Psilocybe montana</i>	·	10	·	10	3.3	10	13.3	1	1
<i>Galerina pumila</i>	4	5	·	9	3.0	4	13.3	3	2
<i>Clitocybe odora</i> var. <i>alba</i>	·	6	1	7	2.3	6	13.3	2	2
<i>Clitocybe vermicularis</i>	6	·	·	6	2.0	6	13.3	1	1
<i>Hyphodontia aspera</i>	·	6	·	6	2.0	5	13.3	2	1
<i>Lepista nuda</i>	·	2	3	5	1.7	3	13.3	2	2
<i>Mycena flavoalba</i>	·	·	5	5	1.7	5	13.3	1	1
<i>Hygrocybe conica</i> var. <i>conica</i>	·	4	·	4	1.3	4	13.3	1	1
<i>Hygrocybe virginea</i> var. <i>virginea</i>	·	2	2	4	1.3	2	13.3	3	2
<i>Phlebiella sulphurea</i>	·	4	·	4	1.3	3	13.3	2	1
<i>Strobilurus stephanocystis</i>	·	2	2	4	1.3	2	13.3	2	2
<i>Ceraceomyces sublaevis</i>	2	1	·	3	1.0	2	13.3	2	2
<i>Resupinatus trichotis</i>	·	3	·	3	1.0	2	13.3	2	1
<i>Leptosporomyces galzinii</i>	·	·	2	2	0.7	2	13.3	1	1
<i>Mycena supina</i>	·	2	·	2	0.7	1	13.3	2	1
<i>Hymenochaete tabacina</i>	36	·	·	36	12.0	36	6.7	1	1
<i>Clitocybe anisata</i>	17	·	·	17	5.7	17	6.7	1	1
<i>Oligoporus alni</i>	11	4	1	16	5.3	7	6.7	4	3
<i>Collybia cookei</i>	15	·	·	15	5.0	15	6.7	1	1
<i>Byssomerulius corium</i>	12	·	·	12	4.0	12	6.7	1	1
<i>Schizophyllum ampulum</i>	·	12	·	12	4.0	12	6.7	1	1
<i>Marasmius setosus</i>	·	·	10	10	3.3	10	6.7	1	1

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Hyphoderma setigerum</i>	8	·	·	8	2.7	8	6.7	1	1
<i>Mycena zephirus</i>	6	1	·	7	2.3	6	6.7	2	2
<i>Coniophora arida</i>	6	·	·	6	2.0	6	6.7	1	1
<i>Cystoderma amianthinum</i>	·	6	·	6	2.0	6	6.7	1	1
<i>Mycena parabolica</i>	·	5	·	5	1.7	5	6.7	1	1
<i>Phanerochaete sanguinea</i>	·	4	·	4	1.3	3	6.7	2	1
<i>Athelia epiphylla</i>	3	·	·	3	1.0	3	6.7	1	1
<i>Hypholoma fasciculare</i>	·	3	·	3	1.0	3	6.7	1	1
<i>Peniophora cinerea</i>	·	3	·	3	1.0	3	6.7	1	1
<i>Gleoporus taxicola</i>	·	2	·	2	0.7	2	6.7	1	1
<i>Psathyrella tephrophylla</i>	1	1	·	2	0.7	1	6.7	2	2
<i>Rhodocybe caelata</i>	2	·	·	2	0.7	2	6.7	1	1
<i>Agaricus dulcidulus</i>	·	1	·	1	0.3	1	6.7	1	1
<i>Calocera viscosa</i>	·	1	·	1	0.3	1	6.7	1	1
<i>Coprinellus xanthothrix</i>	·	·	1	1	0.3	1	6.7	1	1
<i>Crepidotus cesatii</i>	·	·	1	1	0.3	1	6.7	1	1
<i>Cystodermella granulosa</i>	1	·	·	1	0.3	1	6.7	1	1
<i>Galerina</i> spp.	·	1	·	1	0.3	1	6.7	1	1
<i>Geastrum quadrifidum</i>	1	·	·	1	0.3	1	6.7	1	1
<i>Gymnopus putillus</i>	1	·	·	1	0.3	1	6.7	1	1
<i>Hygrocybe laeta</i>	·	·	1	1	0.3	1	6.7	1	1
<i>Tubulicrinis subulatus</i>	·	1	·	1	0.3	1	6.7	1	1
<i>Tulasnella pallida</i>	·	1	·	1	0.3	1	6.7	1	1
<i>Phellinus punctatus</i>	2	·	1	x	x	2	6.7	2	2

Abbreviations: see Tab. A3.2; number of subplots – 15

Table A3.11  
Macrofungi in plot 16 (phase 8, Jelonka reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Cortinarius trivialis</i>	110	45	335	490	163.3	330	100.0	7	3
<i>Amanita citrina</i>	25	3	105	133	44.3	96	100.0	7	3
<i>Amanita muscaria</i>	9	11	16	36	12.0	15	100.0	4	3
<i>Cortinarius</i> spp. II	6	3	88	97	32.3	88	92.3	3	3
<i>Tricholoma equestre</i>	15	17	32	64	21.3	32	92.3	5	3
<i>Russula fragilis</i>	5	1	57	63	21.0	55	84.6	5	3
<i>Rhodocollybia butyracea</i> f. <i>asema</i>	23	25	6	54	18.0	24	76.9	6	3
<i>Cortinarius</i> spp. I	3	9	48	60	20.0	48	69.2	4	3
<i>Russula clavipes</i>	·	5	13	18	6.0	11	69.2	3	2
<i>Russula galochroa</i>	5	6	7	18	6.0	7	61.5	4	3
<i>Cantharellus cibarius</i>	20	81	36	137	45.7	63	53.8	11	3
<i>Inocybe flocculosa</i>	·	·	69	69	23.0	61	53.8	2	1
<i>Lactarius torminosus</i>	8	·	15	23	7.7	15	53.8	3	2
<i>Cortinarius alboviolaceus</i>	3	1	18	22	7.3	18	46.2	3	3
<i>Inocybe</i> cf. <i>margaritispora</i>	1	·	17	18	6.0	17	46.2	2	2
<i>Cortinarius</i> spp. III	·	1	52	53	17.7	52	30.8	2	2
<i>Leccinum scabrum</i>	·	4	2	6	2.0	4	30.8	2	2
<i>Clitopilus prunulus</i>	5	·	15	20	6.7	12	23.1	3	2
<i>Laccaria proxima</i>	2	·	7	9	3.0	6	23.1	3	2
<i>Lactarius uvidus</i>	·	·	8	8	2.7	8	23.1	1	1
<i>Thelephora caryophyllea</i>	·	·	10	10	3.3	7	15.4	2	1
<i>Russula paludosa</i>	3	·	2	5	1.7	2	15.4	3	2
<i>Tricholoma portentosum</i>	3	2	·	5	1.7	3	15.4	2	2

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Tricholoma albobrunneum</i>	·	·	4	4	1.3	4	15.4	1	1
<i>Boletus edulis</i>	1	·	2	3	1.0	1	15.4	3	2
<i>Chalciporus piperatus</i>	·	1	2	3	1.0	1	15.4	3	2
<i>Ramaria cf. abietina</i>	2	·	1	3	1.0	2	15.4	2	2
<i>Hebeloma</i> sp. 1	·	3	·	3	1.0	3	7.7	1	1
<i>Russula mustelina</i>	2	1	·	3	1.0	1	7.7	3	2
<i>Hebeloma</i> sp. 2	·	2	·	2	0.7	2	7.7	1	1
<i>Inocybe lacera</i>	·	·	2	2	0.7	2	7.7	1	1
<i>Paxillus involutus</i>	·	2	·	2	0.7	2	7.7	1	1
<i>Amanita phalloides</i>	·	·	1	1	0.3	1	7.7	1	1
<i>Coltricia perennis</i>	·	1	·	1	0.3	1	7.7	1	1
<i>Hebeloma stenocystis</i>	·	·	1	1	0.3	1	7.7	1	1
<i>Russula versicolor</i>	1	·	·	1	0.3	1	7.7	1	1
<b>Other species</b>									
<i>Clitocybe anisata</i>	61	·	12	73	24.3	59	92.3	3	2
<i>Galerina hypnorum</i>	36	4	38	78	26.0	31	84.6	7	3
<i>Mycena pura</i>	44	2	2	48	16.0	44	84.6	4	3
<i>Gymnopus aquosus</i>	105	15	1	121	40.3	105	76.9	3	3
<i>Clitocybe fragrans</i>	16	·	25	41	13.7	16	76.9	3	2
<i>Mycena leptocephala</i>	19	·	14	33	11.0	19	76.9	2	2
<i>Inonotus rheades</i>	19	8	6	x	x	19	76.9	7	3
<i>Mycena flavoalba</i>	60	6	58	124	41.3	54	69.2	6	3
<i>Lycoperdon molle</i>	26	3	3	32	10.7	20	69.2	7	3
<i>Hypholoma fasciculare</i>	6	14	83	103	34.3	70	61.5	5	3
<i>Mycena olivaceomarginata</i>	2	·	31	33	11.0	31	61.5	2	2
<i>Clitocybe candicans</i>	32	·	6	38	12.7	27	53.8	4	2
<i>Coprinellus xanthothrix</i>	1	1	8	10	3.3	8	53.8	3	3
<i>Tubaria furfuracea</i>	21	27	120	168	56.0	91	46.2	5	3
<i>Flammulaster rhombisporus</i>	12	·	62	74	24.7	62	46.2	3	2
<i>Macrolepiota procera</i>	6	3	4	13	4.3	6	46.2	4	3
<i>Mycena stipitata</i>	7	·	3	10	3.3	7	46.2	3	2
<i>Gymnopus confluens</i>	57	·	3	60	20.0	41	38.5	3	2
<i>Dacrymyces variisporus</i>	·	10	47	57	19.0	47	38.5	2	2
<i>Mycena sanguinolenta</i>	2	·	23	25	8.3	23	38.5	2	2
<i>Galerina</i> spp.	2	·	20	22	7.3	19	38.5	3	2
<i>Conocybe subpubescens</i>	·	6	3	9	3.0	4	38.5	3	2
<i>Coprinellus micaceus</i>	2	15	223	240	80.0	223	30.8	3	3
<i>Collybia cirrhata</i>	·	·	144	144	48.0	144	30.8	1	1
<i>Steccherinum fimbriatum</i>	8	6	36	50	16.7	22	30.8	4	3
<i>Rhodocybe fallax</i>	13	·	27	40	13.3	27	30.8	2	2
<i>Phaeomarasmium erinaceus</i>	2	·	11	13	4.3	11	30.8	2	2
<i>Clitocybe vibecina</i>	·	6	·	6	2.0	6	30.8	1	1
<i>Crepidotus lundellii</i>	186	·	12	198	66.0	126	23.1	3	2
<i>Exidia plana</i>	45	60	14	119	39.7	60	23.1	3	3
<i>Hymenochaete tabacina</i>	·	5	28	33	11.0	18	23.1	4	2
<i>Clitocybe diatreta</i>	13	·	8	21	7.0	13	23.1	2	2
<i>Ceraceomyces crispatus</i>	14	2	·	16	5.3	10	23.1	3	2
<i>Psathyrella candolleana</i>	1	12	2	15	5.0	12	23.1	3	3
<i>Gymnopus androsaceus</i>	8	·	1	9	3.0	8	23.1	2	2
<i>Gymnopus impudicus</i>	1	2	5	8	2.7	5	23.1	3	3
<i>Clitocybe metachroa</i>	7	·	·	7	2.3	7	23.1	1	1
<i>Leptosporomyces galzinii</i>	·	·	5	5	1.7	5	23.1	1	1
<i>Coprinellus disseminatus</i>	140	·	256	396	132.0	200	15.4	3	2
<i>Dacrymyces stillatus</i>	·	200	·	200	66.7	200	15.4	1	1



Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Peniophora polygonia</i>	157	.	1	158	52.7	100	15.4	4	2
<i>Resupinatus trichotis</i>	20	40	.	60	20.0	40	15.4	2	2
<i>Gymnopus acervatus</i>	.	11	23	34	11.3	23	15.4	2	2
<i>Pleurotus ostreatus</i>	.	32	.	32	10.7	20	15.4	2	1
<i>Schizophyllum amplum</i>	4	.	25	29	9.7	25	15.4	2	2
<i>Hypholoma lateritium</i>	9	2	13	24	8.0	13	15.4	3	3
<i>Crucibulum laeve</i>	9	.	8	17	5.7	9	15.4	2	2
<i>Hyphoderma setigerum</i>	7	.	9	16	5.3	9	15.4	2	2
<i>Calocera cornea</i>	14	.	.	14	4.7	14	15.4	1	1
<i>Galerina pumila</i>	12	.	.	12	4.0	12	15.4	1	1
<i>Marasmiellus vaillantii</i>	.	.	8	8	2.7	8	15.4	1	1
<i>Clitocybe phyllophila</i>	7	.	.	7	2.3	7	15.4	1	1
<i>Hygrocybe conica</i> var. <i>conica</i>	.	.	7	7	2.3	7	15.4	1	1
<i>Mycena inclinata</i>	2	2	.	4	1.3	2	15.4	2	2
<i>Hygrocybe virginea</i> var. <i>virginea</i>	1	.	2	3	1.0	2	15.4	2	2
<i>Tremella mesenterica</i>	2	1	.	3	1.0	2	15.4	2	2
<i>Galerina marginata</i>	1	.	1	2	0.7	1	15.4	2	2
<i>Mycena aetites</i>	2	.	.	2	0.7	1	15.4	2	1
<i>Mycena galopus</i>	1	.	1	2	0.7	1	15.4	2	2
<i>Daedaleopsis confragosa</i>	4	.	1	x	x	4	15.4	2	2
<i>Fomes fomentarius</i>	4	3	2	x	x	4	15.4	7	3
<i>Schizopora paradoxa</i>	60	.	.	60	20.0	60	7.7	1	1
<i>Trametes ochracea</i>	32	.	.	32	10.7	17	7.7	2	1
<i>Sphaerobolus stellatus</i>	.	.	30	30	10.0	30	7.7	1	1
<i>Phlebia rufa</i>	.	.	25	25	8.3	25	7.7	1	1
<i>Tubaria confragosa</i>	.	.	13	13	4.3	13	7.7	1	1
<i>Ampulloclitocybe clavipes</i>	.	12	.	12	4.0	12	7.7	1	1
<i>Psathyrella tephrophylla</i>	.	.	10	10	3.3	10	7.7	1	1
<i>Mycena filopes</i>	9	.	.	9	3.0	9	7.7	1	1
<i>Panus suavisissimus</i>	3	.	5	8	2.7	5	7.7	3	2
<i>Mycena flavescens</i>	.	.	6	6	2.0	6	7.7	1	1
<i>Rhodocollybia maculata</i>	.	6	.	6	2.0	6	7.7	1	1
<i>Gymnopus peronatus</i>	.	.	5	5	1.7	5	7.7	1	1
<i>Mycena stylobates</i>	.	.	5	5	1.7	5	7.7	1	1
<i>Mycena vitilis</i>	.	.	5	5	1.7	5	7.7	1	1
<i>Amylostereum laevigatum</i>	4	.	.	4	1.3	4	7.7	1	1
<i>Byssomerulius corium</i>	4	.	.	4	1.3	4	7.7	1	1
<i>Stereum hirsutum</i>	.	.	4	4	1.3	4	7.7	1	1
<i>Hyphodontia quercina</i>	.	.	3	3	1.0	3	7.7	1	1
<i>Calocera viscosa</i>	.	2	.	2	0.7	2	7.7	1	1
<i>Clitocybe nebularis</i>	.	.	2	2	0.7	2	7.7	1	1
<i>Cystoderma amianthinum</i>	.	2	.	2	0.7	2	7.7	1	1
<i>Paneolus papilionaceus</i>	.	2	.	2	0.7	2	7.7	1	1
<i>Chlorophyllum rachodes</i>	.	.	1	1	0.3	1	7.7	1	1
<i>Entoloma</i> spp.	.	.	1	1	0.3	1	7.7	1	1
<i>Geastrum quadrifidum</i>	1	.	.	1	0.3	1	7.7	1	1
<i>Gymnopus ocior</i>	.	1	.	1	0.3	1	7.7	1	1
<i>Hypochnicium detriticum</i>	.	.	1	1	0.3	1	7.7	1	1
<i>Lepiota magnispora</i>	.	1	.	1	0.3	1	7.7	1	1
<i>Lycoperdon nigrescens</i>	.	.	1	1	0.3	1	7.7	1	1
<i>Lycoperdon perlatum</i>	1	.	.	1	0.3	1	7.7	1	1
<i>Mycena metata</i>	.	1	.	1	0.3	1	7.7	1	1
<i>Mycena rubromarginata</i>	1	.	.	1	0.3	1	7.7	1	1
<i>Phellinus punctatus</i>	1	.	.	x	x	1	7.7	1	1

Abbreviations: see Tab. A3.2; number of subplots – 13

Table A3.12  
Macrofungi in plot 2C (phase 9/10, Czechy Orlańskie reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Russula emetica</i>	·	45	2	47	15.7	44	100	3	2
<i>Lactarius rufus</i>	·	28	1	29	9.7	28	100	2	2
<i>Paxillus involutus</i>	3	4	18	25	8.3	18	100	3	3
<i>Xerocomus badius</i>	3	11	6	20	6.7	11	100	4	3
<i>Amanita citrina</i>	·	24	·	24	8.0	24	75	1	1
<i>Cortinarius</i> spp. I	4	20	·	24	8.0	20	75	2	2
<i>Cortinarius</i> spp. II	·	8	9	17	5.7	9	75	2	2
<i>Russula decolorans</i>	·	8	5	13	4.3	8	75	3	2
<i>Russula paludosa</i>	·	6	4	10	3.3	4	75	4	2
<i>Russula fragilis</i>	·	14	·	14	4.7	14	50	1	1
<i>Cortinarius caperatus</i>	·	13	·	13	4.3	13	50	1	1
<i>Thelephora terrestris</i>	·	1	2	3	1.0	2	50	2	2
<i>Russula badia</i>	·	4	·	4	1.3	4	25	1	1
<i>Amanita porphyria</i>	·	3	·	3	1.0	3	25	1	1
<i>Cortinarius semisanguineus</i>	·	3	·	3	1.0	3	25	1	1
<i>Cantharellus cibarius</i>	·	1	·	1	0.3	1	25	1	1
<b>Other species</b>									
<i>Exobasidium vaccinii</i>	2	89	47	138	46.0	80	100	7	3
<i>Gymnopus androsaceus</i>	16	31	42	89	29.7	21	100	8	3
<i>Leptosporomyces galzinii</i>	·	35	·	35	11.7	35	100	1	1
<i>Galerina pumila</i>	5	12	4	21	7.0	9	100	6	3
<i>Mycena zephirus</i>	8	6	·	14	4.7	8	100	3	2
<i>Dacrymyces stillatus</i>	140	210	500	850	283.3	320	75	7	3
<i>Collybia cirrhata</i>	45	·	180	225	75.0	180	75	2	2
<i>Galerina</i> spp.	2	17	·	19	6.3	17	75	2	2
<i>Mycena metata</i>	6	8	·	14	4.7	7	75	4	2
<i>Botryobasidium vagum</i>	·	5	2	7	2.3	4	75	3	2
<i>Entoloma</i> spp.	·	2	3	5	1.7	3	75	2	2
<i>Phlebiella sulphurea</i>	1	2	1	4	1.3	1	75	4	3
<i>Hypholoma capnoides</i>	4	40	·	44	14.7	22	50	4	2
<i>Clitocybe vibecina</i>	·	20	1	21	7.0	18	50	3	2
<i>Mycena galopus</i>	4	2	10	16	5.3	10	50	4	3
<i>Cystoderma amianthinum</i>	5	·	5	10	3.3	5	50	3	2
<i>Mycena purpureofusca</i>	4	1	·	5	1.7	4	50	2	2
<i>Auriscalpium vulgare</i>	·	4	·	4	1.3	3	50	2	1
<i>Gymnopilus bellulus</i>	·	4	·	4	1.3	2	50	2	1
<i>Coniophora arida</i>	·	2	·	2	0.7	2	50	1	1
<i>Gleoporus taxicola</i>	10	·	·	10	3.3	10	25	1	1
<i>Phlebiella pseudotsugae</i>	·	9	·	9	3.0	9	25	1	1
<i>Crepidotus appianatus</i>	·	5	·	5	1.7	5	25	1	1
<i>Gymnopilus penetrans</i>	2	·	3	5	1.7	3	25	2	2
<i>Baeospora myosura</i>	1	1	1	3	1.0	1	25	3	3
<i>Ceraceomyces sublaevis</i>	·	3	·	3	1.0	3	25	1	1
<i>Ampulloclitocybe clavipes</i>	2	·	·	2	0.7	2	25	1	1
<i>Collybia tuberosa</i>	2	·	·	2	0.7	2	25	1	1
<i>Galerina calyptrata</i>	·	2	·	2	0.7	2	25	1	1
<i>Pseudohydnum gelatinosum</i>	2	·	·	2	0.7	2	25	1	1
<i>Tapinella panuoides</i>	2	·	·	2	0.7	2	25	1	1
<i>Clitocybe anisata</i>	·	1	·	1	0.3	1	25	1	1
<i>Hyphodontia aspera</i>	·	1	·	1	0.3	1	25	1	1

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Phanerochaete sanguinea</i>	·	1	·	1	0.3	1	25	1	1
<i>Psilocybe montana</i>	1	·	·	1	0.3	1	25	1	1
<i>Tubulicrinis subulatus</i>	·	·	1	1	0.3	1	25	1	1
<i>Xeromphalina cornui</i>	1	·	·	1	0.3	1	25	1	1

AC – abundance of carpophores during the indicated year; AC3y – abundance of carpophores during 3 years of observations; aACy – average abundance of carpophores per year; mDCv – maximum abundance of carpophores per visit; SF% – spatial frequency of a species (percentage of subplots with the species presence; plot surface – 400 m<sup>2</sup>, number of subplots – 4); tTF – number of visits with a species presence; TFy – number of years with a species presence; x – perennial carpophores

Table A3.13  
Macrofungi in plot 5C (phase 9/10, Czechy Orlańskie reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Paxillus involutus</i>	8	12	23	43	14.3	19	100	5	3
<i>Amanita citrina</i>	·	34	1	35	11.7	29	100	3	2
<i>Russula fragilis</i>	·	20	1	21	7.0	20	75	2	2
<i>Russula vinosa</i>	·	9	1	10	3.3	5	75	3	2
<i>Cantharellus cibarius</i>	1	1	3	5	1.7	1	75	5	3
<i>Cortinarius caperatus</i>	·	5	·	5	1.7	5	75	1	1
<i>Russula emetica</i>	·	5	·	5	1.7	5	75	1	1
<i>Lactarius tabidus</i>	5	7	·	12	4.0	7	50	2	2
<i>Cortinarius vibratilis</i>	·	2	2	4	1.3	2	50	2	2
<i>Laccaria proxima</i>	1	3	·	4	1.3	3	50	2	2
<i>Russula xerampelina</i>	·	4	·	4	1.3	4	50	1	1
<i>Lactarius rufus</i>	1	2	·	3	1.0	2	50	2	2
<i>Russula paludosa</i>	1	1	1	3	1.0	1	50	3	3
<i>Xerocomus badius</i>	1	2	·	3	1.0	2	50	2	2
<i>Russula decolorans</i>	·	1	1	2	0.7	1	50	2	2
<i>Thelephora terrestris</i>	1	6	·	7	2.3	6	25	2	2
<i>Russula badia</i>	1	5	·	6	2.0	4	25	3	2
<i>Cortinarius</i> spp. I	·	2	·	2	0.7	2	25	1	1
<i>Tricholoma saponaceum</i> var. <i>squamosum</i>	·	·	2	2	0.7	2	25	1	1
<i>Amanita porphyria</i>	·	1	·	1	0.3	1	25	1	1
<i>Chroogomphus rutilus</i>	·	1	·	1	0.3	1	25	1	1
<i>Cortinarius</i> spp. II	·	1	·	1	0.3	1	25	1	1
<i>Cortinarius</i> spp. III	·	1	·	1	0.3	1	25	1	1
<i>Rhodocollybia butyracea</i> f. <i>asema</i>	·	·	1	1	0.3	1	25	1	1
<b>Other species</b>									
<i>Dacrymyces stillatus</i>	·	100	301	401	133.7	200	100	5	2
<i>Collybia cirrhata</i>	70	75	70	215	71.7	75	100	3	3
<i>Gymnopus androsaceus</i>	68	35	80	183	61.0	65	100	7	3
<i>Mycena zephirus</i>	108	59	3	170	56.7	94	100	7	3
<i>Hypholoma capnoides</i>	5	29	5	39	13.0	29	100	3	3
<i>Leptosporomyces galzinii</i>	·	24	·	24	8.0	24	100	1	1
<i>Mycena metata</i>	5	2	2	9	3.0	3	100	4	3
<i>Galerina pumila</i>	1	5	1	7	2.3	3	100	5	3
<i>Mycena galopus</i>	4	23	7	34	11.3	21	75	5	3
<i>Exobasidium vaccinii</i>	1	3	14	18	6.0	10	75	6	3
<i>Armillaria ostoyae</i>	·	6	·	6	2.0	6	75	1	1
<i>Hygrophoropsis aurantiaca</i>	2	1	·	3	1.0	2	75	2	2

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Clitocybe vibecina</i>	1	47	·	48	16	47	50	2	2
<i>Mycena vulgaris</i>	16	·	29	45	15.0	29	50	2	2
<i>Collybia tuberosa</i>	9	30	·	39	13.0	30	50	2	2
<i>Xeromphalina cornui</i>	3	·	1	4	1.3	3	50	2	2
<i>Cystoderma amianthinum</i>	1	1	1	3	1.0	1	50	3	3
<i>Phanerochaete sanguinea</i>	·	2	·	2	0.7	1	50	2	1
<i>Panellus mitis</i>	·	·	35	35	11.7	35	25	1	1
<i>Clitocybe diatreta</i>	·	12	·	12	4.0	12	25	1	1
<i>Dacrymyces variisporus</i>	·	·	12	12	4.0	12	25	1	1
<i>Galerina</i> spp.	3	4	·	7	2.3	4	25	2	2
<i>Collybia cookei</i>	5	·	·	5	1.7	5	25	1	1
<i>Botryobasidium vagum</i>	·	3	·	3	1.0	2	25	2	1
<i>Clitocybe anisata</i>	·	3	·	3	1.0	3	25	1	1
<i>Mycena pura</i>	2	·	1	3	1.0	1	25	3	2
<i>Auriscalpium vulgare</i>	2	·	·	2	0.7	2	25	1	1
<i>Galerina mniophila</i>	·	2	·	2	0.7	2	25	1	1
<i>Phlebiella sulphurea</i>	·	2	·	2	0.7	1	25	2	1
<i>Athelia decipiens</i>	1	·	·	1	0.3	1	25	1	1
<i>Baeospora myosura</i>	·	1	·	1	0.3	1	25	1	1
<i>Clitocybe candicans</i>	1	·	·	1	0.3	1	25	1	1
<i>Contophora arida</i>	·	1	·	1	0.3	1	25	1	1
<i>Cystodermella granulosa</i>	·	1	·	1	0.3	1	25	1	1
<i>Macrolepiota procera</i>	·	·	1	1	0.3	1	25	1	1
<i>Mycena purpureofusca</i>	·	·	1	1	0.3	1	25	1	1
<i>Mycena sanguinolenta</i>	1	·	·	1	0.3	1	25	1	1
<i>Inonotus rheades</i>	·	·	1	x	x	1	25	1	1

Abbreviations: see Tab. A3.12

Table A3.14  
Macrofungi in plot 6C (phase 9/10, Czechy Orlańskie reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Paxillus involutus</i>	6	11	28	45	15.0	24	100	4	3
<i>Cortinarius caperatus</i>	1	22	·	23	7.7	22	100	2	2
<i>Amanita citrina</i>	·	35	·	35	11.7	34	75	2	1
<i>Russula emetica</i>	·	22	·	22	7.3	22	75	1	1
<i>Amanita porphyria</i>	·	5	3	8	2.7	5	75	2	2
<i>Russula badia</i>	7	18	·	25	8.3	14	50	3	2
<i>Russula paludosa</i>	·	2	3	5	1.7	3	50	2	2
<i>Russula decolorans</i>	·	1	2	3	1.0	1	50	3	2
<i>Thelephora terrestris</i>	13	·	·	13	4.3	13	25	1	1
<i>Xerocomus badius</i>	2	1	6	9	3.0	4	25	4	3
<i>Cantharellus cibarius</i>	·	8	·	8	2.7	8	25	1	1
<i>Cortinarius vibratilis</i>	6	2	·	8	2.7	6	25	2	2
<i>Cortinarius semisanguineus</i>	1	4	1	6	2.0	4	25	3	3
<i>Rhodocollybia butyracea</i> f. <i>asema</i>	·	·	5	5	1.7	5	25	1	1
<i>Cortinarius</i> spp. III	2	·	·	2	0.7	2	25	1	1
<i>Cortinarius</i> spp. I	·	1	·	1	0.3	1	25	1	1
<i>Tylopilus felleus</i>	·	1	·	1	0.3	1	25	1	1
<b>Other species</b>									
<i>Dacrymyces stillatus</i>	100	350	695	1145	381.7	645	100	5	3
<i>Gymnopus androsaceus</i>	216	14	102	332	110.7	202	100	11	3

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Exobasidium vaccinii</i>	·	66	31	97	32.3	47	100	5	2
<i>Mycena galopus</i>	31	34	3	68	22.7	32	100	6	3
<i>Mycena metata</i>	18	32	1	51	17.0	22	100	6	3
<i>Cystoderma amianthinum</i>	7	9	14	30	10.0	13	100	5	3
<i>Galerina pumila</i>	4	3	13	20	6.7	11	100	5	3
<i>Mycena zephirus</i>	151	41	1	193	64.3	148	75	5	3
<i>Collybia cirrhata</i>	15	30	4	49	16.3	30	75	3	3
<i>Leptosporomyces galzinii</i>	·	18	·	18	6.0	18	75	1	1
<i>Xeromphalina cornui</i>	14	·	·	14	4.7	14	75	1	1
<i>Ampulloclitocybe clavipes</i>	2	8	·	10	3.3	4	75	3	2
<i>Entoloma</i> spp.	·	1	8	9	3.0	5	75	3	2
<i>Auriscalpium vulgare</i>	1	5	·	6	2.0	5	75	2	2
<i>Collybia tuberosa</i>	2	11	34	47	15.7	24	50	5	3
<i>Mycena vulgaris</i>	10	2	20	32	10.7	20	50	3	3
<i>Pholiota spumosa</i>	13	·	1	14	4.7	13	50	2	2
<i>Hygrophoropsis aurantiaca</i>	5	4	1	10	3.3	4	50	5	3
<i>Baeospora myosura</i>	3	1	3	7	2.3	3	50	4	3
<i>Clitocybe vibecina</i>	·	·	4	4	1.3	2	50	2	1
<i>Hyphodontia subalutacea</i>	·	1	1	2	0.7	1	50	2	2
<i>Dacrymyces ovisporus</i>	80	·	·	80	26.7	80	25	1	1
<i>Mycena purpureofusca</i>	2	6	·	8	2.7	6	25	2	2
<i>Mycena adonis</i>	·	4	·	4	1.3	4	25	1	1
<i>Mycena aetites</i>	·	·	4	4	1.3	4	25	1	1
<i>Phlebiella sulphurea</i>	·	2	·	2	0.7	2	25	1	1
<i>Cystoderma jasonis</i> var. <i>jasonis</i>	·	·	1	1	0.3	1	25	1	1
<i>Galerina calyptrata</i>	1	·	·	1	0.3	1	25	1	1
<i>Galerina mniophila</i>	·	1	·	1	0.3	1	25	1	1
<i>Galerina</i> spp.	·	1	·	1	0.3	1	25	1	1
<i>Lycoperdon nigrescens</i>	·	1	·	1	0.3	1	25	1	1
<i>Mycena flavoalba</i>	1	·	·	1	0.3	1	25	1	1
<i>Strobilurus stephanocystis</i>	·	·	1	1	0.3	1	25	1	1

Abbreviations: see Tab. A3.12

Table A3.15  
Macrofungi in plot 7C (phase 9/10, Czechy Orłañskie reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Cortinarius caperatus</i>	4	44	·	48	16.0	44	100	2	2
<i>Russula decolorans</i>	·	10	13	23	7.7	9	100	6	2
<i>Amanita citrina</i>	·	15	1	16	5.3	13	100	3	2
<i>Cortinarius semisanguineus</i>	9	26	26	61	20.3	26	75	5	3
<i>Russula paludosa</i>	1	4	9	14	4.7	8	75	6	3
<i>Xerocomus badius</i>	2	4	2	8	2.7	3	75	4	3
<i>Russula emetica</i>	·	5	·	5	1.7	5	75	1	1
<i>Thelephora terrestris</i>	·	15	·	15	5.0	15	50	1	1
<i>Lactarius rufus</i>	1	13	·	14	4.7	13	50	2	2
<i>Russula badia</i>	2	·	4	6	2.0	4	50	2	2
<i>Russula turci</i>	·	5	·	5	1.7	5	50	1	1
<i>Amanita porphyria</i>	·	3	·	3	1.0	3	50	1	1
<i>Amanita rubescens</i>	1	1	1	3	1.0	1	50	3	3
<i>Paxillus involutus</i>	·	3	·	3	1.0	3	50	1	1
<i>Tricholoma saponaceum</i> var. <i>squamosum</i>	1	5	4	10	3.3	5	25	3	3

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Cortinarius</i> spp. III	·	4	·	4	1.3	4	25	1	1
<i>Laccaria proxima</i>	·	2	·	2	0.7	2	25	1	1
<i>Russula xerampelina</i>	·	2	·	2	0.7	2	25	1	1
<i>Cantharellus cibarius</i>	·	·	1	1	0.3	1	25	1	1
<i>Cortinarius</i> spp. II	·	1	·	1	0.3	1	25	1	1
<i>Cortinarius vibratilis</i>	·	1	·	1	0.3	1	25	1	1
<i>Suillus variegatus</i>	·	·	1	1	0.3	1	25	1	1
<b>Other species</b>									
<i>Exobasidium vaccinii</i>	5	156	49	210	70.0	150	100	6	3
<i>Gymnopus androsaceus</i>	38	45	89	172	57.3	29	100	12	3
<i>Collybia cirrhata</i>	21	19	40	80	26.7	40	100	4	3
<i>Leptosporomyces galzinii</i>	·	28	·	28	9.3	28	100	1	1
<i>Dacrymyces stillatus</i>	165	290	45	500	166.7	240	75	6	3
<i>Xeromphalina cornui</i>	5	12	·	17	5.7	12	75	2	2
<i>Botryobasidium vagum</i>	3	11	·	14	4.7	6	75	3	2
<i>Baeospora myosura</i>	3	6	2	11	3.7	4	75	6	3
<i>Pseudomerulius aureus</i>	15	65	·	80	26.7	50	50	3	2
<i>Hypholoma capnoides</i>	5	4	13	22	7.3	7	50	5	3
<i>Hygrophoropsis aurantiaca</i>	2	5	·	7	2.3	5	50	2	2
<i>Contiophora arida</i>	3	1	2	6	2.0	3	50	4	3
<i>Cystoderma amianthinum</i>	1	2	·	3	1.0	1	50	3	2
<i>Mycena zephirus</i>	1	1	1	3	1.0	1	50	3	3
<i>Galerina</i> spp.	1	1	·	2	0.7	1	50	2	2
<i>Dacrymyces macnabbii</i>	·	50	·	50	16.7	50	25	1	1
<i>Gymnopus ocior</i>	·	26	·	26	8.7	26	25	1	1
<i>Gleoporus taxicola</i>	·	3	·	3	1.0	3	25	1	1
<i>Entoloma</i> spp.	·	·	2	2	0.7	1	25	2	1
<i>Galerina pumila</i>	1	·	1	2	0.7	1	25	2	2
<i>Mycena metata</i>	1	·	1	2	0.7	1	25	2	2
<i>Mycena vulgaris</i>	·	2	·	2	0.7	2	25	1	1
<i>Phlebiella sulphurea</i>	·	2	·	2	0.7	2	25	1	1
<i>Athelia decipiens</i>	1	·	·	1	0.3	1	25	1	1
<i>Clitocybe diatreta</i>	·	1	·	1	0.3	1	25	1	1
<i>Clitocybe vibecina</i>	·	1	·	1	0.3	1	25	1	1
<i>Mycena sanguinolenta</i>	·	1	·	1	0.3	1	25	1	1

Abbreviations: see Tab. A3.12

Table A3.16  
Macrofungi in plot 8C (phase 9/10, Czechy Orlańskie reserve)

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<b>ECM species</b>									
<i>Cortinarius semisanguineus</i>	8	12	8	28	9.3	12	100	5	3
<i>Russula decolorans</i>	·	7	13	20	6.7	9	100	4	2
<i>Russula paludosa</i>	1	3	3	7	2.3	3	100	4	3
<i>Paxillus involutus</i>	7	6	21	34	11.3	21	75	4	3
<i>Lactarius rufus</i>	·	30	·	30	10.0	30	75	1	1
<i>Russula badia</i>	3	11	·	14	4.7	10	75	3	2
<i>Amanita citrina</i>	·	8	1	9	3.0	7	75	3	2
<i>Thelephora terrestris</i>	30	4	·	34	11.3	30	50	2	2
<i>Xerocomus badius</i>	2	6	4	12	4.0	6	50	3	3
<i>Cortinarius vibratilis</i>	2	2	·	4	1.3	2	50	2	2
<i>Cortinarius caperatus</i>	2	1	·	3	1.0	2	50	2	2

Species	AC '94	AC '95	AC '96	AC3y	aACy	mACv	SF%	tTF	TFy
<i>Laccaria proxima</i>	1	1	·	2	0.7	1	50	2	2
<i>Russula xerampelina</i>	·	2	·	2	0.7	2	50	1	1
<i>Amanita porphyria</i>	·	2	·	2	0.7	2	25	1	1
<i>Amanita rubescens</i>	1	·	·	1	0.3	1	25	1	1
<i>Russula vesca</i>	·	1	·	1	0.3	1	25	1	1
<b>Other species</b>									
<i>Collybia cirrhata</i>	35	38	199	272	90.7	186	100	5	3
<i>Gymnopus androsaceus</i>	49	69	60	178	59.3	69	100	8	3
<i>Exobasidium vaccinii</i>	3	13	26	42	14.0	16	100	8	3
<i>Xeromphalina cornui</i>	13	25	·	38	12.7	25	100	2	2
<i>Leptosporomyces galzinii</i>	·	27	·	27	9.0	27	100	1	1
<i>Mycena metata</i>	6	5	11	22	7.3	11	100	5	3
<i>Cystoderma amianthinum</i>	3	1	8	12	4.0	7	100	5	3
<i>Dacrymyces stillatus</i>	·	1570	700	2270	756.7	1570	75	3	2
<i>Mycena zephrus</i>	15	7	·	22	7.3	15	75	2	2
<i>Mycena galopus</i>	6	8	6	20	6.7	6	75	6	3
<i>Galerina pumila</i>	2	5	6	13	4.3	5	75	4	3
<i>Entoloma</i> spp.	·	·	12	12	4.0	12	75	1	1
<i>Phlebiella sulphurea</i>	4	5	·	9	3.0	4	75	4	2
<i>Hyphodontia aspera</i>	1	6	·	7	2.3	6	75	2	2
<i>Hypholoma fasciculare</i>	183	24	39	246	82.0	177	50	8	3
<i>Auriscalpium vulgare</i>	·	4	·	4	1.3	4	50	1	1
<i>Coniophora arida</i>	3	1	·	4	1.3	3	50	2	2
<i>Cystodermella granulosa</i>	2	·	·	2	0.7	2	50	1	1
<i>Dacrymyces ovisporus</i>	·	6	·	6	2.0	6	25	1	1
<i>Gymnopilus penetrans</i>	·	·	5	5	1.7	5	25	1	1
<i>Galerina</i> spp.	·	3	·	3	1.0	3	25	1	1
<i>Gymnopus peronatus</i>	·	·	3	3	1.0	3	25	1	1
<i>Hyphoderma setigerum</i>	3	·	·	3	1.0	3	25	1	1
<i>Galerina mniophila</i>	·	·	2	2	0.7	2	25	1	1
<i>Mycena vulgaris</i>	·	2	·	2	0.7	2	25	1	1
<i>Ampulloclitocybe clavipes</i>	1	·	·	1	0.3	1	25	1	1
<i>Athelia decipiens</i>	·	·	1	1	0.3	1	25	1	1
<i>Baeospora myosura</i>	·	·	1	1	0.3	1	25	1	1
<i>Hyphodontia subalutacea</i>	1	·	·	1	0.3	1	25	1	1
<i>Macrolepiota permixta</i>	·	·	1	1	0.3	1	25	1	1
<i>Phanerochaete sanguinea</i>	1	·	·	1	0.3	1	25	1	1

Abbreviations: see Tab. A3.12





## Appendix 4

### Results of k-mean analysis of fungi

Table A4.1  
Members of clusters and distances from respective cluster centres – ECM species

Species	Distance	Species	Distance
Cluster 1 (8 cases)			
<i>Amanita muscaria</i>	0.789689	<i>Hygrophorus hypothejus</i>	0.746702
<i>Chalciporus piperatus</i>	1.179500	<i>Inocybe lacera</i>	1.497421
<i>Cortinarius cinnamomeoluteus</i>	1.006049	<i>Laccaria proxima</i>	0.991403
<i>Cortinarius</i> spp. I	1.569105	<i>Suillus luteus</i>	0.624135
Cluster 2 (12 cases)			
<i>Boletus edulis</i>	1.204770	<i>Rhodocollybia butyracea</i> f. <i>asema</i>	1.198975
<i>Cortinarius parevernus</i>	1.180209	<i>Thelephora terrestris</i>	1.987908
<i>Cortinarius</i> spp. II	1.860347	<i>Tricholoma albobrunneum</i>	1.154798
<i>Cortinarius</i> spp. III	1.793187	<i>Tricholoma equestre</i>	1.073890
<i>Laccaria bicolor</i>	1.279095	<i>Tricholoma portentosum</i>	0.914971
<i>Lactarius rufus</i>	1.957838	<i>Tricholoma saponaceum</i>	1.162415
Cluster 3 (13 cases)			
<i>Cantharellus cibarius</i>	1.357769	<i>Lactarius uvidus</i>	0.454643
<i>Clitopilus prunulus</i>	1.256667	<i>Leccinum aurantiacum</i>	1.351530
<i>Cortinarius alboviolaceus</i>	0.652723	<i>Russula clavipes</i>	0.405515
<i>Cortinarius trivialis</i>	1.173932	<i>Russula galochroa</i>	0.493669
<i>Hebeloma</i> sp. 3	1.038246	<i>Russula fragilis</i>	1.873435
<i>Inocybe flocculosa</i>	0.879006	<i>Thelephora caryophyllea</i>	0.478748
<i>Lactarius torminosus</i>	0.562072		
Cluster 4 (11 cases)			
<i>Amanita citrina</i>	1.641824	<i>Russula badia</i>	0.456178
<i>Amanita porphyria</i>	0.649528	<i>Russula decolorans</i>	0.613345
<i>Cortinarius caperatus</i>	0.587840	<i>Russula emetica</i>	0.978808
<i>Cortinarius semisanguineus</i>	0.966275	<i>Russula paludosa</i>	0.459364
<i>Cortinarius vibratilis</i>	1.006671	<i>Xerocomus badius</i>	0.649507
<i>Paxillus involutus</i>	1.564870		
Cluster 5 (54 cases)			
<i>Amanita gemmata</i>	0.523716	<i>Inocybe praetervisa</i>	0.493738
<i>Amanita pantherina</i>	0.746301	<i>Lactarius necator</i>	0.444144
<i>Amanita phalloides</i>	0.330851	<i>Lactarius quieticolor</i>	0.777556
<i>Amanita rubescens</i>	1.243976	<i>Lactarius tabidus</i>	0.884765
<i>Boletus pinophilus</i>	0.287742	<i>Lactarius vietus</i>	0.334876
<i>Boletus reticulatus</i>	0.379404	<i>Leccinum scabrum</i>	0.550001
<i>Chroogomphus rutilus</i>	0.408213	<i>Ramaria</i> cf. <i>abietina</i>	0.428299
<i>Coltricia perennis</i>	0.389802	<i>Ramaria eumorpha</i>	1.282514
<i>Cortinarius acetosus</i>	0.285980	<i>Rhizopogon luteolus</i>	0.340235
<i>Cortinarius anthracinus</i>	1.252337	<i>Russula</i> cf. <i>mustelina</i>	0.428299
<i>Cortinarius cinnabarinus</i>	0.474943	<i>Russula subfetens</i>	0.340537
<i>Cortinarius colus</i>	0.974179	<i>Russula turci</i>	0.692938
<i>Cortinarius mucosus</i>	1.214468	<i>Russula versicolor</i>	0.330851

Species	Distance	Species	Distance
<i>Cortinarius multififormis</i> var. <i>coniferarum</i>	0.750410	<i>Russula vesca</i>	0.436899
<i>Cortinarius purpurascens</i>	0.908160	<i>Russula vinosa</i>	0.842382
<i>Cortinarius sanguineus</i>	0.515556	<i>Russula vinosobrunnea</i>	0.321510
<i>Cortinarius</i> spp. IV	0.655574	<i>Russula xerampelina</i>	0.878182
<i>Gyroporus castaneus</i>	0.462315	<i>Sarcodon squamosus</i>	1.036126
<i>Hebeloma stenocystis</i>	0.330851	<i>Scleroderma fuscum</i>	0.413826
<i>Hebeloma</i> sp. 1	0.769771	<i>Suillus bovinus</i>	0.398441
<i>Hebeloma</i> sp. 2	0.637234	<i>Suillus variegatus</i>	0.412319
<i>Hygrophorus hypothejus</i> var. <i>aureus</i>	0.455719	<i>Tricholoma imbricatum</i>	0.812475
<i>Inocybe assimilata</i>	0.890619	<i>Tricholoma pessundatum</i>	0.381157
<i>Inocybe asterospora</i>	1.104966	<i>Tricholoma saponaceum</i> var. <i>squamosum</i>	1.176912
<i>Inocybe calida</i>	0.639305	<i>Tylopilus felleus</i>	0.447547
<i>Inocybe curvipes</i>	0.944775	<i>Xerocomus chrysenteron</i>	0.367833
<i>Inocybe</i> cf. <i>margaritispota</i>	0.763932	<i>Xerocomus subtomentosus</i>	0.598075

Table A4.2  
Members of clusters and distances from respective cluster centres – non-ECM species

Species	Distance	Species	Distance
Cluster 1 (6 cases)			
<i>Clitocybe diatreta</i>	1.890329	<i>Entoloma</i> spp.	1.462573
<i>Collybia cirrhata</i>	1.745173	<i>Galerina</i> spp.	1.331584
<i>Cystoderma amianthinum</i>	0.932323	<i>Gymnopus androsaceus</i>	1.764856
Cluster 2 (11 cases)			
<i>Ampulloclitocybe clavipes</i>	1.619621	<i>Mycena flavoalba</i>	1.533014
<i>Cantharellula umbonata</i>	2.187800	<i>Mycena leptocephala</i>	1.407885
<i>Clavaria argillacea</i>	1.912443	<i>Mycena olivaceomarginata</i>	1.737105
<i>Galerina atkinsoniana</i>	1.158011	<i>Psilocybe montana</i>	1.832456
<i>Hygroaster asterosporus</i>	1.185667	<i>Rickenella fibula</i>	0.950151
<i>Lycoperdon perlatum</i>	1.239052		
Cluster 3 (47 cases)			
<i>Amylostereum laevigatum</i>	0.557839	<i>Hyphoderma setigerum</i>	0.655903
<i>Byssomerulius corium</i>	0.514562	<i>Hypholoma fasciculare</i>	1.811642
<i>Clitocybe anisata</i>	0.886725	<i>Inonotus rheades</i>	0.415238
<i>Clitocybe candicans</i>	0.985087	<i>Lycoperdon molle</i>	1.161743
<i>Clitocybe fragrans</i>	1.245170	<i>Macrolepiota procera</i>	0.915513
<i>Clitocybe metachroa</i>	0.397062	<i>Marasmiellus vaillantii</i>	0.858866
<i>Clitocybe phyllophila</i>	0.928363	<i>Marasmius epiphyllus</i>	0.823209
<i>Coprinellus disseminatus</i>	1.176315	<i>Mycena filopes</i>	0.358211
<i>Coprinellus micaceus</i>	1.101572	<i>Mycena pura</i>	1.177489
<i>Crepidotus lundellii</i>	0.872516	<i>Mycena sanguinolenta</i>	0.472491
<i>Crucibulum laeve</i>	0.312945	<i>Mycena stipata</i>	0.353784
<i>Cystoderma carcharias</i>	0.829006	<i>Mycena stylobates</i>	0.721379
<i>Dacrymyces variisporus</i>	1.770178	<i>Mycena vitilis</i>	1.117191
<i>Daedaleopsis confragosa</i>	0.484802	<i>Peniophora polygonia</i>	0.792184
<i>Exidia plana</i>	0.546914	<i>Phaeomarasmium erinaceus</i>	0.338036
<i>Flammulaster granulosus</i>	1.237628	<i>Phlebia rufa</i>	0.317687
<i>Flammulaster rhombosporus</i>	0.508864	<i>Psathyrella candolleana</i>	0.408357
<i>Galerina hypnorum</i>	1.629354	<i>Resupinatus trichotis</i>	0.657706
<i>Galerina marginata</i>	0.572542	<i>Schizophyllum amplum</i>	0.390806
<i>Gleocystidiellum porosum</i>	0.824749	<i>Sphaerobolus stellatus</i>	0.607043
<i>Gymnopus aquosus</i>	0.668549	<i>Stereum hirsutum</i>	0.545113

Species	Distance	Species	Distance
<i>Gymnopus confluens</i>	1.125006	<i>Tremella mesenterica</i>	0.608938
<i>Hygrocybe conica</i>	0.774268	<i>Tubaria furfuracea</i>	0.831217
<i>Hymenochaete tabacina</i>	0.340679		
Cluster 4 (14 cases)			
<i>Auriscalpium vulgare</i>	1.177597	<i>Leptosporomyces galzinii</i>	1.125963
<i>Baeospora myosura</i>	1.149912	<i>Mycena galopus</i>	1.036916
<i>Collybia tuberosa</i>	1.581279	<i>Mycena metata</i>	0.773333
<i>Dacrymyces stillatus</i>	3.121296	<i>Mycena vulgaris</i>	1.472430
<i>Exobasidium vaccini</i>	1.376177	<i>Mycena zephyrus</i>	0.940386
<i>Galerina pumila</i>	1.529068	<i>Phlebiella sulphurea</i>	0.970005
<i>Hypholoma capnoides</i>	1.579716	<i>Xeromphalina cornui</i>	1.053344
Cluster 5 (124 cases)			
<i>Agaricus dulcidulus</i>	0.265466	<i>Lepiota pseudolilacea</i>	0.347784
<i>Armillaria ostoyae</i>	0.702082	<i>Lepista nuda</i>	0.411007
<i>Athelia decipiens</i>	0.531347	<i>Lycoperdon dermoxanthum</i>	1.307389
<i>Athelia epiphylla</i>	0.336987	<i>Lycoperdon ericaeum</i>	0.899553
<i>Botryobasidium vagum</i>	1.206346	<i>Lycoperdon excipuliforme</i>	0.474054
<i>Calocera cornea</i>	0.590135	<i>Lycoperdon marginatum</i>	0.675753
<i>Calocera furcata</i>	0.473336	<i>Lycoperdon nigrescens</i>	1.001204
<i>Calocera viscosa</i>	0.254998	<i>Lycoperdon norvegicum</i>	0.423829
<i>Ceraceomyces crispatus</i>	0.618744	<i>Macrolepiota mastoidea</i>	0.368881
<i>Ceraceomyces sublaevis</i>	0.573990	<i>Macrolepiota permixta</i>	0.387989
<i>Chlorophyllum rhacodes</i>	0.246993	<i>Marasmius oreades</i>	1.597007
<i>Clitocybe brumalis</i>	0.309632	<i>Marasmius setosus</i>	0.538533
<i>Clitocybe nebularis</i>	0.279329	<i>Melanoleuca graminicola</i>	0.688612
<i>Clitocybe odora</i>	0.785122	<i>Mycena adonis</i>	0.895200
<i>Clitocybe odora</i> var. <i>alba</i>	0.469758	<i>Mycena aetites</i>	0.633033
<i>Clitocybe vermicularis</i>	0.442004	<i>Mycena arcangeliana</i>	0.296801
<i>Clitocybe vibecina</i>	1.139557	<i>Mycena flavescens</i>	0.424183
<i>Collybia cookei</i>	1.160069	<i>Mycena inclinata</i>	0.358777
<i>Coniophora arida</i>	1.057121	<i>Mycena parabolica</i>	0.458667
<i>Conocybe ochrostriata</i>	0.336987	<i>Mycena purpureofusca</i>	0.985813
<i>Conocybe rickeniana</i>	0.302984	<i>Mycena rubromarginata</i>	0.387850
<i>Conocybe spiculoides</i>	0.309632	<i>Mycena septentrionalis</i>	0.288917
<i>Conocybe subpubescens</i>	0.499668	<i>Mycena speirea</i>	1.121998
<i>Coprinellus xanthothrix</i>	0.507932	<i>Mycena supina</i>	0.294516
<i>Coprinus friesii</i>	0.887246	<i>Mycetinis scorodonius</i>	1.560248
<i>Crepidotus applanatus</i>	0.646016	<i>Oligoporus alni</i>	0.636647
<i>Crepidotus cesatii</i>	0.257884	<i>Panellus mitis</i>	1.179276
<i>Crepidotus variabilis</i>	0.999158	<i>Paneolus pabillonaceus</i>	0.885524
<i>Crinipellis scabella</i>	1.838806	<i>Panus suavissimus</i>	0.476879
<i>Cystoderma jasonis</i>	0.858008	<i>Parasola plicatilis</i>	0.625791
<i>Cystoderma jasonis</i> var. <i>lilacipes</i>	1.099015	<i>Peniophora cinerea</i>	0.336987
<i>Cystodermella granulosa</i>	1.050270	<i>Phanerochaete sanguinea</i>	0.726263
<i>Dacrymyces macnabbii</i>	1.187326	<i>Phellinus punctatus</i>	0.308319
<i>Dacrymyces ovisporus</i>	1.447944	<i>Phlebiella pseudotsugae</i>	0.773731
<i>Entoloma conferendum</i>	0.673462	<i>Pholiota gumosa</i>	0.558960
<i>Entoloma jubatum</i>	1.223281	<i>Pholiota spumosa</i>	0.879249
<i>Entoloma nitens</i>	0.296801	<i>Pleurotus ostreatus</i>	0.773829
<i>Entoloma rhodopolium</i>	0.356133	<i>Polyporus ciliatus</i>	0.320874
<i>Entoloma undatum</i>	0.315975	<i>Psathyrella fibrillosa</i>	0.303412
<i>Fomes fomentarius</i>	0.499668	<i>Psathyrella gyroflexa</i>	0.835502
<i>Galerina badipes</i>	0.296357	<i>Psathyrella tephrophylla</i>	0.527476
<i>Galerina calyptrata</i>	0.530320	<i>Pseudohydnum gelatinosum</i>	0.470388

Species	Distance	Species	Distance
<i>Galerina mniophila</i>	0.913767	<i>Pseudomerulius aureus</i>	1.301152
<i>Geastrum quadrididum</i>	0.219100	<i>Resinicium bicolor</i>	0.246993
<i>Gleoporus taxicola</i>	0.933332	<i>Rhodocollybia maculata</i>	0.424183
<i>Gymnopilus bellulus</i>	0.600051	<i>Rhodocybe caelata</i>	0.787312
<i>Gymnopilus penetrans</i>	0.885718	<i>Rhodocybe fallax</i>	0.825488
<i>Gymnopilus acervatus</i>	0.787797	<i>Rhodocybe hirneola</i>	0.654539
<i>Gymnopus dryophilus</i>	0.964499	<i>Rhodocybe parilis</i>	1.400197
<i>Gymnopus impudicus</i>	0.814622	<i>Ripartites tricholoma</i>	0.624283
<i>Gymnopus ocior</i>	1.021795	<i>Schizopora paradoxa</i>	0.920818
<i>Gymnopus peronatus</i>	0.628279	<i>Steccherinum fimbriatum</i>	0.877747
<i>Gymnopus putillus</i>	0.545022	<i>Strobilurus stephanocystis</i>	1.574688
<i>Hygrocybe laeta</i>	0.257884	<i>Strobilurus tenacellus</i>	0.494816
<i>Hygrocybe virginea</i> var. <i>virginea</i>	1.266556	<i>Tapinella atrotomentosa</i>	0.334966
<i>Hygrophoropsis aurantiaca</i>	1.179235	<i>Tapinella panuoides</i>	0.470388
<i>Hyphoderma praetermissum</i>	0.769009	<i>Trametes ochracea</i>	0.773829
<i>Hyphodontia aspera</i>	0.849322	<i>Trichaptum fuscoviolaceum</i>	0.902484
<i>Hyphodontia quercina</i>	0.320255	<i>Tubaria confragosa</i>	0.574484
<i>Hyphodontia subalutacea</i>	0.544677	<i>Tubulicrinis subulatus</i>	0.351309
<i>Hypholoma lateritium</i>	0.751848	<i>Tulasnella pallida</i>	0.257884
<i>Lepiota magnispora</i>	0.246993	<i>Vesiculomyces citrinus</i>	0.685440

## Appendix 5

### List of Basidiomycota in the old-field successional series near the Białowieża old-growth forest (Jelonka reserve and Czechy Orlańskie reserve) with notes on distribution and substrates

J – Jelonka reserve; CzO – Czechy Orlańskie reserve; I–XII – months; 1, 11, 67, 85 – plot numbers; E, V, R, I – The Red Data Book categories according to WOJEWODA & ŁAWRYNOWICZ (2006).

#### BASIDIOMYCOTA EXOBASIDIOMYCETES

##### Exobasidiales

*Exobasidium vaccinii* (Fuckel) Woronin – on leaves of *Vaccinium vitis-idaea*; CzO: 2, 5, 6, 7, 8, phase 9/10; X 1994, VI–X 1995, VI–XI 1996, VII 1997.

#### DACRYMYCETES

##### Dacrymycetales

*Calocera cornea* (Batsch) Fr. – on log of birch; J: 16, phase 8, X 1994.

*Calocera furcata* (Fr.) Fr. – on log of pine; J: 7, phase 6, X 1994; R.

*Calocera viscosa* (Pers.) Fr. – at a rotting stump; J: 11, 16, phase 8, IX 1995, VIII 1996.

*Dacrymyces macnabbii* D.A. Reid – on fallen branch of pine; CzO: 7, phase 9/10, VI 1995; rev. W. Wojewoda.

*Dacrymyces ovisporus* Bref. – on fallen branches of pine; CzO: 6, 8, phase 9/10, IX 1994, IX 1995; E; rev. W. Wojewoda.

*Dacrymyces stillatus* Nees – on fallen branches and logs of pine; J: 84, 1, 11, 16, phases 4, 6/7, 8, IX–X 1994, VI, X–XI 1995, VI, IX, XI 1996; CzO: 2, 5, 6, 7, 8, phase 9/10, VIII–X 1994, VI, VIII–X 1995, V–VI, X 1996.

*Dacrymyces variisporus* McNabb – on dead and living branches of pine; J: 7, 13, 1, 6, 11, 16, phases 6–8, IX–X 1994, VI, IX, XI 1995, VI, IX, XI 1996; CzO: 5, phase 9/10, IX 1996; E.

#### TREMELLOMYCETES

##### Tremellales

*Exidia plana* Donk – on branches of aspen; J: 11, 16, phase 8; IX–X 1994, VI 1995, VI, XI 1996.

*Pseudohydnum gelatinosum* (Scop.) P. Karst. – on rotten stump; CzO: 2, 5, phase 9/10; IX 1994, IX 1997.

*Tremella mesenterica* Retz. – on twigs of aspen; J: 7, 11, 16, phases 6, 8; X 1994, VI 1995, VI–VIII 1996.

#### AGARICOMYCETES

##### Agaricales

*Agaricus dulcidulus* Schulzer [= *Agaricus semotus* Fr.] – on soil; J: 1, 11, phases 6/7, 8; IX 1994, X 1995.

*Amanita citrina* Pers. – on soil; J: 84, 11, 16, phases 4, 8; CzO: 2, 5, 6, 7, 8, phase 9/10; IX–X 1994, VIII–X 1995, VIII–X 1996.

*Amanita gemmata* (Fr.) Bertill. – on soil; J: 11, phase 8; IX 1994, VI 1995.

*Amanita muscaria* (L.) Lam. – on soil; J: 67, 84, 61, 85, 7, 13, 1, 6, 11, 16, phases 4–8; IX 1994, IX–XI 1995, VII–XI 1996, VII 1998.

*Amanita pantherina* (DC.) Krombh. – on soil; J: 84, 85, 7, 13, 6, 11, phases 4–8; IX 1994, VI–VIII 1995, X 1995, VII 1996.

*Amanita phalloides* (Vaill. ex Fr.) Link – on soil; J: 16, phase 8; VIII 1995, IX 1998.

*Amanita porphyria* Alb. & Schwein. – on soil, under pine; CzO: 2, 5, 6, 7, 8, phase 9/10; X 1995, VIII 1996, VII 1997, VII 1998.

*Amanita rubescens* (Pers.) Gray – on soil; J: 84, 85, 7, 1, 6, 11, phases 4–8; VI 1994, VIII, X 1995, VII–IX 1996; CzO: 5, 7, 8, phase 9/10; IX 1994, X 1995, VIII 1996, VII 1997.

*Ampulloclitocybe clavipes* (Pers.) Redhead, Lutzoni, Moncalvo & Vilgalys [= *Clitocybe clavipes* (Pers.) P. Kumm.] – on soil and on litter; J: 67, 84, 61, 7, 13, 1, 6, 11, 16, phases 4-8, IX-X 1994, IX-XI 1995, VIII-XI 1996; CzO: 2, 6, 8, phase 9/10, IX 1994, IX-X 1995, IX 1998.

*Armillaria ostoyae* (Romagn.) Herink – on stump; CzO: 5, phase 9/10; X 1995.

*Baeospora myosura* (Fr.) Singer – on buried cones of pine; J: 7, 1, 6, 11, phases 6-8; IX-X 1994, IX-XI 1995, XI 1996, XI 1997, CzO: 6, 7, 8, phase 9/10, IX-X 1994, IX-XI 1995, X-XI 1996.

*Cantharellula umbonata* (J.F. Gmel.) Singer – on mosses – *Polytrichum juniperinum* and *P. commune*; J: 67, 85, 7, phases 4-6; IX-X 1994, IX-XI 1995, VII-XI 1996, VII 1998.

*Ceraceomyces crispatus* (O.F. Müll.) Rauschert [= *Ceraceomyces serpens* (Tode) Ginns] – on fallen twigs of aspen; J: 16, phase 8, VI, X 1994, XI 1996; E; *det. E. Parmasto*.

*Ceraceomyces sublaevis* (Bres.) Jülich – on fallen branches of aspen, pine; J: 11, phase 8, X 1994, X 1995; CzO: 2, phase 9/10, X 1995; *det. W. Wojewoda*.

*Chlorophyllum rachodes* (Vittad.) Vellinga [= *Macrolepiota rhacodes* ('rachodes') (Vittad.) Singer] – on soil; J: 16, phase 8, X 1995.

*Clavaria argillacea* Pers. – on soil, under *Calluna vulgaris*; J: 67, 84, 61, 85, 7, 13, 1, 6, phases 4-6/7, X 1994, X-XI 1995, X-XI 1996; R.

*Clitocybe anisata* Velen. – on soil and on litter; J: 1, 11, 16, phases 6/7, 8; IX-X 1994, IX 1995; CzO: 2, 5, phase 9/10, X 1995; *det. H. Komorowska*.

*Clitocybe brumalis* (Fr.) Gillet – among grass and herbs; J: 82, phase 2, X 1995; *det. H. Komorowska*.

*Clitocybe candicans* (Pers.) P. Kumm. – on soil and on litter; J: 7, 13, 6, 11, 16, phases 6-8, IX-X 1994, IX-X 1995, X 1996; CzO: 5, phase 9/10, IX 1994; *det. H. Komorowska*.

*Clitocybe diatrete* (Fr.) P. Kumm. – on soil and on litter, most frequently under *Juniperus*; J: 67, 84, 61, 85, 7, 13, 1, 6, 11, 16, phases 4-8, VI, IX-X 1994, VIII-XI 1995, VII-XI 1996, VII, IX-X 1997, VII 1998; CzO: 5, 7, phase 9/10, X 1995; *det. H. Komorowska*.

*Clitocybe fragrans* (With.) P. Kumm. – on soil and on litter; J: 61, 7, 1, 6, 11, 16, phases 5-8, X 1994, IX-X 1995, X-XI 1996, X 1997; *det. H. Komorowska*.

*Clitocybe metachroa* (Fr.) P. Kumm. – on litter; J: 11, 16, phase 8, X 1994, X 1995, X 1996; *det. H. Komorowska*.

*Clitocybe nebularis* (Batsch) P. Kumm. – on soil; J: 16, phase 8, X 1995.

*Clitocybe odora* (Bull.) P. Kumm. var. *odora* – on soil and on litter; J: 85, 11, phases 5, 8, IX 1994, X-XI 1995; *det. H. Komorowska*.

*Clitocybe odora* var. *alba* J.E. Lange – on soil and on litter; J: 11, phase 8, X 1995, VII 1996; *det. H. Komorowska*.

*Clitocybe phyllophila* (Pers.) P. Kumm. – on litter; J: 1, 11, 16, phases 6/7-8, IX 1994, IX-X 1995, X 1996; *det. H. Komorowska*.

*Clitocybe vermicularis* (Fr.) Quél. – on litter; J: 11, phase 8, VI 1994; *det. H. Komorowska*.

*Clitocybe vibecina* (Fr.) Quél. [= *Clitocybe langei* Singer ex Hora] – on soil and on litter; J: 84, 85, 7, 13, 1, 6, 16, phases 4-8, X 1994, IX-XI 1995, X-XI 1996, X-XI 1997; CzO: 2, 5, 6, 7, phase 9/10, IX 1994, X-XI 1995, X-XI 1996, X 1997; *det. H. Komorowska*.

*Clitopilus prunulus* (Scop.) P. Kumm. – on soil; J: 67, 84, 85, 7, 13, 1, 6, 11, 16, phases 4-8, IX 1994, IX-X 1995, VII-VIII, X 1996.

*Collybia cirrhata* (Pers.) Quél. – on decaying sporocarps of fungi; J: 67, 84, 61, 85, 7, 13, 1, 6, 11, 16, phases 4-8, IX-X 1994, IX-XI 1995, IX-XI 1996; CzO: 2, 5, 6, 7, 8, phase 9/10, IX 1994, IX 1995, IX-XI 1996.

*Collybia cookei* (Bres.) J.D. Arnold – on decaying sporocarps of fungi; J: 61, 85, 11, phases 5, 8, IX-X 1994, IX 1995, VIII 1996; CzO: 5, phase 9/10, IX 1994.

*Collybia tuberosa* (Bull.) P. Kumm. – on decaying sporocarps of fungi; J: 85, 7, 13, 1, phases 5-6/7, IX-X 1994, IX-X 1995, IX-X 1996; CzO: 2, 5, 6, phase 9/10, IX-X 1994, IX-X 1995, X-XI 1996.

*Conocybe ochrostriata* Hauskn. [= *Conocybe siennophylla* (Berk. & Broome) Singer s. auct.] – on humus; J: 11, phase 8, X 1995.

*Conocybe pulchella* (Velen.) Hauskn. & Švrček [= *Conocybe pseudopilosella* Kühner & Watling] – on litter; CzO, phase 9/10, XI 1997.

*Conocybe rickeniana* P.D. Orton – on soil, among mosses; J: 84, phase 4, VI 1994.

*Conocybe spiculoides* Kühner & Watling. – among grass and herbs; J: 82, phase 2, X 1995.

*Conocybe subpubescens* P.D. Orton [= *Conocybe cryptocystis* (G.F. Akt.) Singer s. auct.] – on humus; J: 16, phase 8, IX 1995, IX, XI 1996.

*Coprinellus disseminatus* (Pers.) J.E. Lange [= *Coprinus disseminatus* (Pers.) Gray] – on rotting wood; J: 16, phase 8, IX 1994, IX-X 1995, VII 1997.



*Coprinellus micaceus* (Bull.) Vilgalys, Hopple & Jacq. Johnson [= *Coprinus micaceus* (Bul.) Fr.] – on rotting wood, on humus; J: 16, phase 8, X 1994, X 1995, VIII 1996.

*Coprinellus xanthothrix* (Romagn.) Vilgalys, Hopple & Jacq. Johnson [= *Coprinus xanthothrix* Romagn.] – on rotting wood, on humus; J: 11, 16, phase 8, IX 1994, IX 1995, IX 1996, VII, IX 1997.

*Coprinopsis friesii* (Quél.) P. Karst. [= *Coprinus friesii* Quél.] – on dead grass; J: 82, phase 2, VII-VIII 1996.

*Cortinarius acetosus* (Velen.) Melot [= *Cortinarius rigens* (Pers.) Fr.] – on soil, under pine; J: 13, phase 6, X 1995; E.

*Cortinarius alboviolaceus* (Pers.) Fr. – on soil, under birch, aspen; J: 11, 16, phase 8, IX 1994, IX-X 1995, VIII, X 1996.

*Cortinarius anthracinus* Fr. – on soil, under pine, among mosses and lichens; J: 61, 13, 1, 6, phases 5-6/7, IX-X 1994, X 1995, VIII, X 1996, IX-X 1998; V.

*Cortinarius caperatus* (Pers.) Fr. [= *Rozites caperatus* (Pers.) P. Karst.] – on soil, under pine; CzO: 2, 5, 6, 7, 8, phase 9/10, IX 1994, X 1995, VII 1998.

*Cortinarius cinnabarinus* Fr. – on soil, under pine; J: 61, 13, phases 5, 6, X 1995, XI 1996.

*Cortinarius cinnamomeoluteus* P.D. Orton – on soil, under pine; J: 67, 84, 61, 85, 7, 13, 1, 6, 11, phases 4-8, VI, IX-X 1994, IX-X 1995, VII-XI 1996, VII, XI 1997, IX 1998.

*Cortinarius colus* Fr. [= *Cortinarius miniatopus* J.E. Lange] – on soil, under pine; J: 7, phase 6, X 1995.

*Cortinarius fulvescens* Fr. – on soil, under pine, among mosses; J: 85, phase 5, X 1998; E.

*Cortinarius mucosus* (Bull.) J.J. Kickx – on soil, under pine; J: 7, 13, 1, 6, phases 6-6/7, IX-X 1994, IX-X 1995, VIII, X 1996.

*Cortinarius multififormis* f. *coniferarum* (M.M. Moser) Nespiak – on soil, under pine; J: 13, 1, 6, phases 6-6/7, X 1994, X 1995, IX-X 1996; R.

NOTE. The specimens match the description of the form proposed by NESPIAK (1975).

*Cortinarius obtusus* (Fr.) Fr. s.l. – on soil, under pine, among mosses and lichens; J: 84, 1, phases 4, 6/7, IX 1998.

*Cortinarius orellanus* Fr. – on soil, under pine; CzO: phase 9/10, VII 1997; R.

*Cortinarius parevernus* Rob. Henry – on soil, under pine; J: 7, 13, 1, 6, 11, phases 6-8, X 1994, X 1995, X-XI 1996, X 1998.

*Cortinarius phoeniceus* (Vent.) Maire [= *Cortinarius purpureus* (Pers.) Fuckel s. auct.] – on soil, under pine, among mosses; J: 1, phase 6/7, VII 1997.

*Cortinarius purpurascens* (Fr.) Fr. – on soil, under pine; J: 67, 7, 13, 1, 6, phases 4, 6, 6/7, X 1994, X 1995, X-XI 1996.

*Cortinarius sanguineus* (Wulfen) Gray – on soil, under pine, among mosses and lichens; J: 1, phase 6/7, X 1995.

*Cortinarius semisanguineus* (Fr.) Fr. – on soil, under pine; J: 1, phase 7, VII 1997; CzO: 2, 6, 6/7, 8, phase 9/10, IX-X 1994, X 1995, X-XI 1996, VII, X 1997, VII 1998.

*Cortinarius trivialis* J.E. Lange s.l. – on soil, under aspen and pine; J: 11, 16, phase 8, IX-X 1994, IX-X 1995, VII-VIII, X 1996, XI 1997, VII 1998.

*Cortinarius vibratilis* (Fr.) Fr. – on soil, under pine; CzO: 5, 6, 7, 8, phase 9/10, IX 1994, X 1995, X 1996.

*Crepidotus applanatus* (Pers.) P. Kumm. – on fallen twigs; CzO: 2, phase 9/10, IX 1995; R.

*Crepidotus cesatii* (Rabenh.) Sacc. [= *Crepidotus sphaerosporus* (Pat.) J.E. Lange] – on fallen twigs; J: 11, phase 8, VI 1996.

*Crepidotus lundellii* Pilát [= *Crepidotus amygdalosporus* Kühner] – on fallen twigs; J: 11, 16, phase 8, IX-X 1994, VI, VIII 1995, VI 1996.

*Crepidotus variabilis* (Pers.) P. Kumm. – on dead stems of *Calluna vulgaris*; J: 84, 61, phases 4-5, IX-X 1995.

*Crinipellis scabella* (Alb. & Schwein.) Murrill [= *Crinipellis stipitaria* (Fr.) Pat.] – among grass and herbs; J: 82, 85, 7, phases 2, 5, 6, IX-X 1994, IX-XI 1995, VII-IX 1996.

*Crucibulum laeve* (Huds.) Kambly – on fallen twigs, on pieces of bark; J: 11, 16, phase 8, IX 1994, IX 1995, VII-VIII 1996.

*Cystoderma amianthinum* (Scop.) Fayod – on soil, among mosses; J: 82, 67, 84, 61, 85, 7, 13, 1, 6, 11, 16, phases 2-8, IX-X 1994, VIII-XI 1995, VIII-XI 1996, X-XI 1997; CzO: 2, 5, 6, 7, 8, phase 9/10, IX-X 1994, X-XI 1995, X-XI 1996, X-XI 1997.

*Cystoderma carcharias* (Pers.) Fayod – on litter; J: 1, 6, 11, phases 6/7-8, X 1994, X 1995, X-XI 1996.

*Cystoderma jasonis* (Cooke & Massee) Harmaja var. *jasonis* [= *C. longisporum* (Kühner) Arnolds] – on soil, among mosses; J: 85, 7, 1, phases 5-6/7, X 1994, X 1995, IX-X 1996, XI 1997.

*Cystoderma jasonis* (Cooke & Massee) Harmaja var. *lilacipes* (Harmaja) I. Saar [= *Cystoderma lilacipes* Harmaja; *C. longisporum* (Kühner) Arnolds var. *purpurascens* Arnolds] – on soil, among mosses; J: 84, 85, 7, 13, 1, phases 4-6/7, X 1994, X 1995, X-XI 1996, XI 1997; CzO, phase 9/10, XI 1997, X 1998.

*Cystoderma granulosa* (Batsch) Harmaja [= *Cystoderma granulorum* (Batsch) Fayod] – on soil, among mosses; J: 85, 7, 13, 1, 11, phases 5-8, IX-X 1994, X 1995, IX-X 1996, XI 1997; CzO: 5, 8 phase 9/10, IX 1994, X 1995; I.

*Entoloma conferendum* (Britzelm.) Noordel. var. *conferendum* – on soil; J: 85, phase 5, X 1996.

*Entoloma jubatum* (Fr.) P. Karst. – on soil; J: 84, 61, 85, 7, 13, phases 4-6, VI 1994, X 1995, XI 1996, IX 1998.

*Entoloma nitens* (Velen.) Noordel. – on soil; J: 13, phase 6, X 1995.

*Entoloma rhodopolium* (Fr.) P. Kumm. var. *rhodopolium* – on soil, among mosses; J: 84, phase 4, X 1995.

*Entoloma undatum* (Fr.) M.M. Moser – on soil; J: 82, 1, phases 2, 6/7, X 1995.

*Flammulaster granulatus* (J.E. Lange) Watling – on fallen twigs, on pieces of bark; J: 84, 85, 7, 6, 11, phases 4-8, IX 1994, IX-X 1995, VII, IX 1996, VII 1997.

*Flammulaster rhombosporus* (G.F. Atk.) Watling – on fallen twigs, on pieces of bark; J: 11, 16, phase 8 IX-X 1994, IX-X 1995, XI 1996; R.

*Galerina atkinsoniana* A.H. Sm. – among mosses; J: 67, 84, 61, 85, 7, 13, 1, 6, phases 4-6/7, IX-X 1994, IX-XI 1995, VII-VIII, X-XI 1996, VII 1997, VII, IX 1998.

*Galerina badipes* (Pers.) Kühner – among mosses; J: 7, phase 6, X 1994; E.

*Galerina calyptata* P.D. Orton – among mosses; J: 13, phase 6, VII 1997; CzO: 2, 6, phase 9/10, IX 1994, X 1995.

*Galerina hypnorum* (Schränk.) Kühner – among mosses; J: 67, 84, 85, 7, 1, 6, 11, 16, phases 4-8, IX-X 1994, IX-XI 1995, VII, X-XI 1996, VII, X-XI 1997; CzO, phase 9/10, XI 1997.

*Galerina marginata* (Batsch) Kühner s.l. [= *Galerina unicolor* (Vahl) Singer] – on fallen twigs; J: 11, 16, phase 8, IX-X 1994, X 1995, XI 1997.

*Galerina mniophila* (Lasch) Kühner – among mosses; J: 1, phase 6/7, VI 1994; CzO: 5, 6, 8, phase 9/10, X 1995, X 1996, XI 1997.

*Galerina pumila* (Pers.) M. Lange – among mosses; J: 67, 84, 61, 85, 7, 13, 1, 11, 16, phases 4-8, IX-X 1994, IX-XI 1995, IX-X 1996, X-XI 1997; CzO: 2, 5, 6, 7, 8, phase 9/10, X 1994, IX-XI 1995, X-XI 1996, X-XI 1997.

*Gymnopilus bellulus* (Peck) Murrill – on fallen branch; CzO: 2, phase 9/10, IX-X 1995.

*Gymnopilus penetrans* (Fr.) Murrill – on fallen branches, on rotting stump; CzO: 2, 8, phase 9/10, X 1994, X 1996.

*Gymnopus acervatus* (Fr.) Murrill [= *Collybia acervata* (Fr.) P. Kumm.] – on decayed wood ; J: 16, phase 8, VI 1995, VI 1996.

*Gymnopus androsaceus* (L.) Antonín & Noordel. [= *Marasmius androsaceus* (L.) Fr.; *Setulipes androsaceus* (L.) Antonín] – on fallen litter of pine, rarely on fallen leaves of aspen; J: 67, 84, 61, 85, 7, 13, 1, 6, 11, 16, phases 4-8, IX-X 1994, VI, IX-XI 1995, VI-VII, IX-XI 1996; CzO: 2, 5, 6, 7, 8, phase 9/10, VIII-X 1994, VI, IX-XI 1995, VI-XI 1996, VII 1997.

*Gymnopus aquosus* (Bull.) Antonín & Noordel. [= *Collybia aquosa* (Bull.) P. Kumm.] – on soil and on litter; J: 67, 84, 61, 7, 11, 16, phases 4-6, 8, VI 1994, VI 1995, VI-VII 1996, VII 1997.

*Gymnopus confluens* (Pers.) Antonín, Halling & Noordel. [= *Collybia confluens* (Pers.) P. Kumm.] – on litter; J: 13, 11, 16, phases 6, 8, IX-X 1994, VIII-XI 1995, VII-IX 1996.

*Gymnopus dryophilus* (Bull.) Murrill [= *Collybia dryophila* (Bull.) P. Kumm.] – on soil and on litter; J: 7, 13, 11, 16, phases 6, 8 IX 1994, X 1995, X 1996, IX 1997.

*Gymnopus impudicus* (Fr.) Antonín, Halling & Noordel. [= *Micromphale impudicum* (Fr.) P.D. Orton] – on soil; J: 85, 1, 16, phases 5, 6/7, 8, IX-X 1994, IX 1995, IX 1996, VII 1998.

*Gymnopus ocior* (Pers.) Antonín & Noordel. [= *Collybia ocior* (Pers.) Vilgalys & O.K. Mill.] – on rotting wood, on litter; J: 16, phase 8, VII 1996; CzO: 7, phase 9/10, VI 1995; E.

*Gymnopus peronatus* (Bolton) Antonín, Halling & Noordel. [= *Collybia peronata* (Bolton) P. Kumm.] – on litter; J: 16, phase 8, IX 1995, VII, X 1997; CzO: 8, phase 9/10, IX 1996.

*Gymnopus putillus* (Fr.) Antonín, Halling & Noordel. [= *Collybia putilla* (Fr.) Singer] – on soil and on litter; J: 84, 61, 7, 13, 1, 6, 11, phases 4-8, X 1994, IX 1995; E.

*Hebeloma stenocystis* J. Favre ex Quadr. – on soil; J: 16, phase 8, X 1995; det. H.J. Beker.

*Hebeloma* sp. 1 – on soil, under willow, aspen, pine; J: 67, 84, 13, 16, phases 4, 6, 8, X 1994, X-XI 1995, X 1996.

*Hebeloma* sp. 2 – on soil; J: 67, 16, phases 4, 8, X 1995, X 1996.

*Hebeloma* sp. 3 – on soil; J: 11, phase 8, IX-X 1994, X 1995, X 1996.

*Hygroaster asterosporus* (J.E. Lange) Singer [= *Omphaliaster asterosporus* (J.E. Lange) Lamoure] – on soil, among mosses and lichens; J: 67, 84, 61, 85, 7, 13, 1, 6, phases 4-6/7, X 1994, X-XI 1995, X-XI 1996, X 1997; E.

*Hygrocybe acutoconica* (Clem.) Singer [= *Hygrocybe persistens* (Britzelm.) Singer] – on soil; J: 82, phase 2, VII 1997; R.

*Hygrocybe conica* (Schaeff.) P. Kumm. var. *conica* – on soil; J: 85, 7, 1, 6, 11, 16, phases 5-8, X 1995.

*Hygrocybe laeta* (Pers.) P. Kumm. – on humus; J: 11, phase 8, XI 1996; V.

*Hygrocybe virginea* var. *virginea* (Wulfen) P.D. Orton & Watling [= *Camarophyllus virgineus* (Wulfen) P. Kumm.] – on soil; J: 82, 84, 61, 85, 13, 1, 11, 16, phases 2-8, X 1994, X 1995, X-XI 1996, XI 1997.

*Hygrophorus aureus* Arrh. [= *Hygrophorus hypothejus* (Fr.) Fr. var. *aureus* (Arrh.) Imler] – on soil, among mosses; J: 67, phase 4, XI 1995; E.

NOTE. According to WOJEWODA & ŁAWRYNOWICZ (1992) the species is threatened and has been assigned to category E. As WOJEWODA (2003) does not treat it as a separate species any more but as a variety of *Hygrophorus hypothejus*, it has not been listed in the latest red list (WOJEWODA & ŁAWRYNOWICZ 2006).

*Hygrophorus hypothejus* (Fr.) Fr. – on soil, most frequently among mosses; J: 67, 84, 61, 85, 7, 13, 1, 6, 11, phases 4-8, X 1994, IX-XI 1995, X-XI 1996, XI 1997; I.

*Hypholoma capnoides* (Fr.) P. Kumm. – on stump; CzO: 2, 5, 7, phase 9/10, X 1994, V, X-XI 1995, X-XI 1996, X 1997.

*Hypholoma fasciculare* (Huds.) P. Kumm. – on decayed wood stake, on stump; J: 84, 7, 6, 11, 16, phases 4, 6-8, IX 1994, VIII, X 1995, X-XI 1996; CzO: 6, 8, phase 9/10, IX-X 1994, VIII-X 1995, VIII, X-XI 1996, IX 1997.

*Hypholoma lateritium* (Schaeff.) P. Kumm. – on rotting wood; J: 1, 6, 16, phases 6/7-8, X 1994, X 1995, X-XI 1996.

*Inocybe assimilata* Britzelm. [= *Inocybe umbrina* Bres.] – on soil; J: 84, 7, 13, 1, 6, phases 4, 6-6/7, X 1994, X-XI 1995, VII-VIII 1996.

*Inocybe asterospora* Quél. – on soil; J: 67, 84, 7, 1, phases 4, 6-6/7, X 1995, VII 1997.

*Inocybe calida* Velen. [= *Inocybe brunneorufa* Stangl & Veselský] – on soil; J: 67, 84, 85, 7, 6, phases 4-6/7, X 1994, X 1995, VII 1997.

*Inocybe curvipes* P. Karst. [= *Inocybe lanuginella* (J. Schröt.) Konrad & Maubl.] – on soil; J: 67, 85, 7, 13, 1, 6, phases 4-6/7, X 1995, VII, XI 1996, VII 1997.

*Inocybe flocculosa* (Berk.) Sacc. [= *Inocybe gausapata* Kühner] – on soil, under birch, aspen; J: 16, phase 8, IX-X 1995, VII 1997.

*Inocybe lacera* (Fr.) P. Kumm. var. *lacera* – on soil, under pine; J: 67, 84, 61, 85, 7, 13, 1, 6, 16, phases 4-8, VI, IX-X 1994, VI, IX-XI 1995, V, VII-XI 1996, XI 1997, IX 1998.

*Inocybe* cf. *margaritispora* (Berk.) Sacc. [= *Inocybe* cf. *phaeosticta* Furrer-Ziogas] – on soil; J: 16, phase 8, IX 1994, X 1995.

*Inocybe praetervisa* Quél. – on soil; J: 84, 61, 85, 13, 6, phases 4-6/7, X 1995, VII, X 1996, VII 1997, IX 1998.

*Laccaria bicolor* (Maire) P.D. Orton – on soil, frequently among mosses; J: 84, 61, 85, 7, 13, 1, 6, phases 4-6/7, IX 1994, VIII-XI 1995, VII-VIII, X-XI 1996, IX 1997.

*Laccaria proxima* (Boud.) Pat. – on soil, frequently among mosses; J: 67, 84, 61, 85, 7, 13, 1, 6, 11, 16, phases 4-8, VI, IX-X 1994, IX-XI 1995, V, VII-XI 1996, XI 1997; CzO: 5, 7, 8, phase 9/10, IX 1994, X-XI 1995, XI 1997.

*Lepiota magnispora* Murrill [= *Lepiota ventriosospora* D.A. Reid] – on soil; J: 16, phase 8, X 1996.

*Lepiota pseudolilacea* Huijsman [= *Lepiota pseudohelveola* Kühner ex Hora] – on soil; J: 13, phase 6, X 1995.

*Lepista nuda* (Bull.) Cooke – on soil and on litter; J: 11, phase 8, X 1995, X 1996, XI 1997.

*Lycoperdon dermoxanthum* Vittad. [= *Bovista dermoxantha* (Vittad.) De Toni; *Bovista pusilla* (Batsch) Pers.] – on soil; J: 67, 84, 61, 85, 7, 13, phases 4-6, IX 1994, IX-X 1995, VIII-IX 1996, VII, X 1997, IX 1998.

*Lycoperdon ericaeum* Bonord. – on soil; J: 85, phase 5, IX-X 1994, IX-X 1995; E.

*Lycoperdon excipuliforme* (Scop.) Pers. [= *Calvatia excipuliformis* (Scop.) Perdeck] – on soil; J: 13, phase 6, IX-X 1994, X 1995.

*Lycoperdon marginatum* Vittad. – on soil; J: 61, 13, phases 5-6, IX 1994, VIII-X 1995, IX, XI 1996, VII 1997; E.

*Lycoperdon molle* Pers. – on soil; J: 67, 84, 85, 13, 11, 16, phases 4-6, 8, VI, IX-X 1994, VIII-XI 1995, VII-XI 1996, VII, IX 1997.

*Lycoperdon nigrescens* Pers. – on soil; J: 67, 84, 85, 11, 16, phases 4, 5, 8, IX-X 1994, IX-X 1995, VIII-IX, XI 1996; CzO: 6, phase 9/10, IX 1995, IX 1997.

*Lycoperdon norvegicum* Demoulin – on soil; J: 84, 85, phases 4-5, X 1994, VIII 1996, VII 1997; I.

*Lycoperdon perlatum* Pers. – on soil; J: 67, 84, 85, 7, 13, 1, 6, 11, 16, phases 4-8, IX-X 1994, IX-XI 1995, X-XI 1996, IX, X 1997, VII 1998.

*Lycoperdon utriforme* Bull. [= *Calvatia utriformis* (Bull.) Jaap] – on soil; J: near the plot no. 61, phase 5, VII 1997.

*Macrolepiota mastoidea* (Fr.) Singer [= *Macrolepiota konradii* (P.D. Orton) M.M. Moser] – on soil; J: 84, 61, phase 4-5, IX 1994, X 1995, IX 1998.

*Macrolepiota permixta* (Barla) Pacioni – on soil; CzO: 8, phase 9/10, XI 1996.

*Macrolepiota procera* (Scop.) Singer – on soil; J: 84, 61, 6, 11, 16, phases 4, 5, 6/7, 8, IX-X 1994, IX-X 1995, VIII, X 1996, IX 1998; CzO: 5, phase 9/10, XI 1996.

*Marasmiellus vaillantii* (Pers.) Singer – on fallen leaves of aspen; J: 85, 7, 11, 16, phases 5-6, 8, IX 1994, VIII-IX 1995, VII, IX 1996, VII 1998; *rev. M. Lisiewska*.

*Marasmius epiphyllus* (Pers.) Fr. – on fallen leaves of aspen; J: 11, phase 8, X 1995, IX 1996; *rev. M. Lisiewska*.

*Marasmius oreades* (Bolton) Fr. – on soil and on litter; J: 82, 61, 13, 1, 11, phases 2, 5-8, IX-X 1994, VI, IX-XI 1995, VI-XI 1996, VII 1996.

*Marasmius setosus* (Sowerby) Noordel. – on fallen leaves of aspen; J: 11, phase 8, IX 1996; V; *rev. M. Lisiewska*.

*Melanoleuca graminicola* (Velen.) Kühner & Maire – on soil; J: 7, 13, phase 6, IX-X 1994, IX-X 1995.

NOTE: acc. to KNUDSEN & VESTERHOLT (2008) the species belongs to the complex of *M. melaleuca* (Pers.: Fr.) Murril s.l.

*Mycena adonis* (Bull.) Gray var. *adonis* – on fallen needles of pine; J: 61, phase 5, IX 1994, IX 1995; CzO: 6, phase 9/10, IX 1995; R; *rev. M. Lisiewska*.

*Mycena aetites* (Fr.) Quél. – on litter; J: 84, 1, 6, 16, phases 4, 6/7, 8, IX-X 1994, IX-X 1995; CzO: 6, phase 9/10, X 1996.

*Mycena arcangeliana* Bres. [= *Mycena lineata* (Fr.) Quél. ss. Lange] – on litter; J: 13, phase 6, XI 1996.

*Mycena epipterygia* (Scop.) Gray var. *epipterygia* – on litter; CzO, phase 9/10, XI 1997.

*Mycena filopes* (Bull.) P. Kumm. – on litter; J: 11, 16, phase 8, IX-X 1994, X 1996.

*Mycena flavescens* Velen. – on litter; J: 16, phase 8, IX 1995; R.

*Mycena flavoalba* (Fr.) Quél. – on decayed herbaceous plant debris, on litter; J: 82, 67, 84, 61, 85, 7, 13, 1, 6, 11, 16, phases 2-8, IX-X 1994, IX-X 1995, IX-XI 1996, XI 1997; CzO: 6, phase 9/10, IX 1994.

*Mycena galopus* (Pers.) P. Kumm. – on litter; J: 85, 13, 11, 16, phases 5, 6, 8, IX 1994, IX-X 1995, VII 1997; CzO: 2, 5, 6, 8, phase 9/10, IX-X 1994, IX-XI 1995, X-XI 1996, X 1997.

*Mycena inclinata* (Fr.) Quél. – on rotting stump; J: 16, phase 8, IX 1994, X 1996.

*Mycena leptcephala* (Pers.) Millet – on decayed herbaceous plant debris, on litter; J: 82, 67, 84, 61, 85, 7, 13, 1, 6, 11, 16, phases 2-8, IX-X 1994, VI, IX-X 1995, VI, IX-X 1996, XI 1997; *rev. M. Lisiewska*.

*Mycena metata* (Fr.) P. Kumm. [= *M. tenella* (Fr.) Quél.] – on litter, on fallen needles of pine; J: 84, 7, 6, 16, phases 4, 6, 6/7, 8, X 1994, XI 1995, XI 1996, CzO: 2, 5, 6, 7, 8, phase 9/10, IX-X 1994, IX-XI 1995, X-XI 1996, XI 1997.

*Mycena olivaceomarginata* (Masse) Masse [= *M. avenacea* (Fr.) Quél.; *M. plicosa* (Fr.) P. Kumm.] – on decayed herbaceous plant debris, on litter; J: 82, 67, 84, 61, 85, 13, 1, 11, 16, phases 2-8, IX-X 1994, VI, IX-XI 1995, IX 1996; *rev. M. Lisiewska*.

*Mycena parabolica* (Fr.) Quél. – on rotting wood; J: 13, 11, phases 6, 8, X 1994, X 1995; *rev. M. Lisiewska*.

*Mycena pura* (Pers.) P. Kumm. – on humus, on litter; J: 82, 67, 84, 61, 7, 13, 1, 6, 11, 16, phases 2-8, 6, VI, IX 1994, IX-X 1995, VII-XI 1996, VII 1997; CzO: 5, phase 9/10, IX-X 1994, XI 1996.

*Mycena purpureofusca* (Peck) Sacc. – on rotting wood; CzO: 2, 5, 6, phase 9/10, IX 1994, IX 1995, X 1996; V.

*Mycena rubromarginata* (Fr.) P. Kumm. – on fallen twigs and decayed wood; J: 11, 16, phase 8, IX-X 1994, XI 1997.

*Mycena sanguinolenta* (Alb. & Schwein.) P. Kumm. – on litter; J: 7, 11, 16, phases 6, 8, IX 1994, IX 1995; CzO: 5, 7, phase 9/10, IX 1994, IX 1995.

*Mycena septentrionalis* Maas Geest. [= *Mycena vitrea* (Fr.) Quél.] – on litter, among mosses; J: 1, phase 6/7, X 1995.

*Mycena smithiana* Kühner [= *M. debilis* (Fr.) Quél.] – on fallen needles of *Juniperus*; J: 16, phase 8, X 1997; *det. M. Lisiewska*.

*Mycena speirea* (Fr.) Gillet – on fallen twigs; J: 13, 1, 6, phases 6, 6/7, X 1994, IX 1996, XI 1997.

*Mycena stipata* Maas Geest. & Schwöbel [= *M. alcalina* (Fr.) P. Kumm. s. auct. p.p.] – on rotting wood; J: 11, 16, phase 8, VI, IX 1994, IX-X 1995.

*Mycena stylobates* (Pers.) P. Kumm. – on fallen twigs; J: 84, 61, 85, 7, 1, 6, 11, 16, phases 4-8, IX 1994, IX-X 1995, VI, IX 1996.

*Mycena supina* (Fr.) P. Kumm. – on stump; J: 11, phase 8, IX-X 1995; E; *det. M. Lisiewska*.

*Mycena vitilis* (Fr.) Quél. – on litter, on fallen twigs; J: 67, 85, 7, 13, 1, 6, 11, 16, phases 4-8, IX-X 1994, IX-XI 1995, VII, X 1996.

*Mycena vulgaris* (Pers.) P. Kumm. – on fallen needles of pine; J: 7, phase 6, X 1994, X 1996; CzO: 5, 6, 7, 8, phase 9/10, X 1994, XI 1995, X 1996.

*Mycena zephrus* (Fr.) P. Kumm. – on litter; J: 11, phase 8, IX 1994, X 1995. CzO: 2, 5, 6, 7, 8, phase 9/10, IX-X 1994, IX-XI 1995, X-XI 1996, X 1997.

*Mycetinis scorodonius* (Fr.) Wilson & Desjardin [= *Marasmius scorodonius* (Fr.) Fr.] – among grass and herbs; J: 67, 84, phase 4, IX-X 1994, IX-XI 1995, VII-IX 1996.

*Panaeolus papilionaceus* (Bull.) Quél. – on hare dung; J: 67, 85, 6, 16, phases 4, 5, 6/7, 8, IX-X 1995, X-XI 1996, VII 1997; R.

*Panellus mitis* (Pers.) Singer – on fallen branches of pine; J: 1, phase 6/7, XI 1995; CzO: 5, phase 9/10, X 1996.

*Parasola plicatilis* (Curtis) Redhead, Vilgalys & Hopple [= *Coprinus plicatilis* (Curtis) Fr.] – among grass and herbs; J: 82, phase 2, VI 1995, VI 1996.

*Phaeomarasmius erinaceus* (Fr.) Kühner – on fallen twigs; J: 13, 11, 16, phases 6, 8, X 1994, VI 1995, VI 1996, VII 1997; R.

*Pholiota gummosa* (Lasch) Singer – on rotting wood; J: 7, phase 6, X 1995, XI 1996.

*Pholiota spumosa* (Fr.) Singer – on rotting stump; CzO: 6, phase 9/10, IX 1994, X 1996, XI 1997.

*Pleurotus ostreatus* (Jacq.) P. Kumm. – on dead trunk of aspen; J: 16, phase 8, X-XI 1996.

*Psathyrella candolleana* (Fr.) Maire – on humus and litter; J: 11, 16, phase 8, VI, IX 1994, V, IX-X 1995, VII-VIII 1996.

*Psathyrella fibrillosa* (Pers.) Maire – on soil; J: 13, 1, phases 6-6/7, X 1995.

*Psathyrella gyroflexa* (Fr.) ss. M.M. Moser – on soil; J: 67, 61, phases 4-5, IX-X 1995.

*Psathyrella tephrophylla* (Romagn.) M.M. Moser ex Bon – on humus; J: 11, 16, phase 8, IX 1994, IX-X 1995.

*Pseudomerulius aureus* (Fr.) Jülich – on fallen branch of pine; CzO: 7, phase 9/10, IX 1994, VI, X 1995; R; *det. E. Parmasto*.

*Psilocybe montana* (Pers.) P. Kumm. – on soil, among mosses and lichens; J: 67, 84, 61, 85, 7, 13, 1, 6, 11, phases 4-8, VI, IX-X 1994, VI, VIII-XI 1995, V-XI 1996; CzO: 2, phase 9/10, IX 1994; R.

*Psilocybe phyllogena* (Sacc.) Peck [= *Psilocybe rhombispora* (Britzelm.) Sacc.] – on soil, among mosses and lichens; J: 67, 7, 13, 1, 6, phases 4, 6-6/7, VII, IX 1997; R.

*Resupinatus trichotis* (Pers.) Singer – on fallen branches; J: 11, 16, phase 8, X 1994, VI, X 1995, VI 1996.

*Rhodocollybia butyracea* (Bull.) Lennox f. *asema* (Fr.) Antonín, Halling & Noordel. [= *Collybia butyracea* (Bull.) P. Kumm. var. *asema* (Fr.) Quél.] – on soil and litter; J: 67, 84, 61, 85, 7, 13, 1, 6, 11, 16, phases 4-8, IX-X 1994, X-XI 1995, VII, X-XI 1996, XI 1997, IX 1998; CzO: 5, 6, phase 9/10, X 1996.

*Rhodocollybia maculata* (Alb. & Schwein.) Singer [= *Collybia maculata* (Alb. & Schwein.) P. Kumm.] – on litter; J: 16, phase 8, VI 1996.

*Rhodocybe caelata* (Fr.) Maire – on soil, among mosses and lichens; J: 67, 85, 7, 13, 6, 11, phases 4-8, IX-X 1994, IX 1995, IX 1996; E.

*Rhodocybe fallax* (Quél.) Singer – on humus; J: 16, phase 8, IX 1994, IX 1995; E.

*Rhodocybe hirneola* (Fr.) P.D. Orton – on soil, among mosses and lichens; J: 67, 85, 13, phases 4-6, X 1994, X 1995, VII 1998.

*Rhodocybe parilis* (Fr.) Singer – on soil, among mosses and lichens; J: 84, 61, 85, 7, 13, 6, phases 4-6/7, IX-X 1994, IX-X 1995, VII, IX, XI 1996, VII 1998.

*Ripartites tricholoma* (Alb. & Schwein.) P. Karst. s.l. – on soil; J: 7, 13, 1, phases 6-6/7, phases 4-6/7 IX-X 1994, X 1995.

*Schizophyllum amplum* (Lév.) Nakasone 191996 [= *Auriculariopsis ampla* (Lév.) Maire] – on dead branches of aspen; J: 11, 16, phase 8, X 1994, VI 1995.

*Strobilurus stephanocystis* (Hora) Singer – on buried pine cones; J: 7, 13, 1, 6, 11, phases 6-8, VI 1994, V 1995, V-VI 1996; CzO: 6, phase 9/10, VI 1996.

*Strobilurus tenacellus* (Pers.) Singer – on buried pine cones; J: 11, phase 8, VI 1994, V 1995.

*Stropharia semiglobata* (Batsch) Quél. – on litter; J: 61, phase 5, X 1997.

*Tapinella atrotomentosa* (Batsch) Šutara [= *Paxillus atrotomentosus* (Batsch) Fr.] – on decayed wood stakes; J: 85, 61, 7, 6, phases 5-6/7, VII-VIII 1996, VII 1997.

*Tapinella panuoides* (Fr.) E.-J. Gilbert [= *Paxillus panuoides* (Fr.) Fr.] – on fallen branch of pine; CzO: 2, phase 9/10, IX 1994.

*Tricholoma albobrunneum* (Pers.) P. Kumm. – on soil, under pine; J: 84, 61, 85, 7, 13, 1, 6, 16, phases 4-8, X 1994, X-XI 1995, X-XI 1996, X 1997, IX 1998; *det. H. Komorowska*.

*Tricholoma equestre* (L.) P. Kumm. – on soil, under pine; J: 61, 85, 7, 13, 1, 6, 11, 16, phases 5-8, IX-X 1994, X-XI 1995, VIII-XI 1996; I.

*Tricholoma imbricatum* (Fr.) P. Kumm. – on soil, under pine; J: 7, 13, 11, phases 6, 8, X 1995, X 1996, X 1998; *det. H. Komorowska*.

*Tricholoma pessundatum* (Fr.) Quél. – on soil, under pine; J: 7, phase 6, X 1994; R; *det. H. Komorowska*.

*Tricholoma portentosum* (Fr.) Quél. – on soil, under pine; J: 84, 61, 7, 13, 1, 6, 11, 16, phases 4-8, X 1994, X-XI 1995, X-XI 1996.



*Tricholoma saponaceum* (Fr.) P. Kumm. var. *saponaceum* – on soil, under pine; J: 61, 7, 13, 1, 6, phases 5-6/7, X 1994, X 1995, IX-XI 1996, X-XI 1997.

*Tricholoma saponaceum* (Fr.) P. Kumm. var. *squamosum* (Cooke) Rea – on soil, under pine; J: 84, phase 4, X 1995; CzO: 5, 7, phase 9/10, IX 1994, X 1995, XI 1996, IX 1998.

*Tubaria confragosa* (Fr.) Harmaja – on fallen branch; J: 16, phase 8, IX 1995; E.

*Tubaria furfuracea* (Pers.) Gillet – on fallen twigs, on pieces of bark; J: 84, 11, 16, phases 4, 8, IX 1994, IX-X 1995, VI-VII, IX-XI 1996, VII, XI 1997.

*Xeromphalina cornui* (Quél.) J. Favre – on fallen needles of pine; J: 11, phase 8, IX-X 1994, IX 1995; CzO: 2, 5, 6, 7, 8, phase 9/10, IX 1994, IX 1995, VII 1996, VII, IX 1997.

#### Atheliales

*Athelia decipiens* (Höhn. & t Litsch.) J. Erikss. – on fallen twigs of pine; CzO: 5, 7, 8, phase 9/10, VIII 1994, VI 1996.

*Athelia epiphylla* Pers. – on decayed log; J: 11, phase 8, X 1994; *det. W. Wojewoda*.

*Leptosporomyces galzinii* (Bourdot) Jülich – on fallen branches, on fallen cones of pine; J: 7, 13, 1, 6, 11, 16, phases 6-8, VI 1995, VI 1996; CzO: 2, 5, 6, 7, 8, phase 9/10, VI 1995; *det. W. Wojewoda*.

#### Boletales

*Boletus edulis* Bull. – on soil, under pine, birch, aspen; J: 67, 84, 61, 85, 13, 1, 6, 11, 16, phases 4-8, IX-X 1994, VI, VIII, IX-X 1995, VII-X 1996, VII 1997, VII 1998.

*Boletus pinophilus* Pilát & Dermek – on soil, under pine; J: 1, 6, phase 6/7, VIII, X 1996, VII 1997.

*Boletus reticulatus* Schaeff. – on soil, under *Betula*, pine; J: 84, 1, phases 4, 6/7, VIII 1995, VIII 1996.

*Chalciporus piperatus* (Bull.) Bataille – on soil, under pine; J: 67, 84, 61, 85, 7, 13, 1, 6, 11, 16, phases 4-8, IX 1994, VIII-X 1995, VII-XI 1996, VII 1998.

*Chroogomphus rutilus* (Schaeff.) O.K. Mill. – on soil, under pine; CzO: 5, phase 9/10, X 1995.

*Coniophora arida* (Fr.) P. Karst. – on fallen branches of pine; J: 11, phase 8, X 1994; CzO: 2, 5, 7, 8, phase 9/10, IX 1994, X 1995, VI, X 1996; *det. W. Wojewoda*.

*Gyroporus castaneus* (Bull.) Quél. – on soil; J: 84, 85, phases 4-5, VIII, X 1995, VIII 1996, VII 1997; R.

*Gyroporus cyanescens* (Bull.) Quél. – on soil; CzO: phase 9/10, VII 1997; R.

*Hygrophoropsis aurantiaca* (Wulfen) Maire – on soil; J: 6, phase 6/7, IX 1996; CzO: 5, 6, 7, phase 9/10, IX-X 1994, X-XI 1995, XI 1996.

*Leccinum aurantiacum* (Bull.) Gray – on soil, under aspen; J: 84, 85, 7, 1, 11, phases 4-8, IX 1994, VI, VIII-X 1995, VII-IX 1996, VII 1997, VII 1998.

*Leccinum scabrum* (Bull.) Gray – on soil, under birch; J: 67, 84, 16, phases 4, 8, IX 1994, VIII, X 1995, VIII 1996, VII, IX 1997.

*Paxillus involutus* (Batsch) Fr. – on soil, under pine, birch, aspen; J: 67, 84, 61, 85, 7, 13, 1, 11, 16, phases 4-8, X 1994, IX-X 1995, VII-VIII, X-XI 1996, VII, XI 1997, VII 1998; CzO: 2, 5, 6, 7, 8, phase 9/10, IX-X 1994, X-XI 1995, VIII, X-XI 1996, XI 1997, VII 1998.

*Rhizopogon luteolus* Fr. & Nordholm [= *R. obtextus* (Spring.) Rauschert] – hypogeous, partly visible on the soil surface; J: 67, 7, 6, phases 4, 6-6/7, IX 1994, X 1995, VIII 1996, XI 1997, IX 1998; *det. M. Ławrynowicz*.

*Scleroderma bovista* Fr. – on soil; J: near the plot no. 11, phase 8, IX 1997.

*Scleroderma fuscum* (Corda) E. Fischer. – on soil; J: 85, phase 5, X 1995; E.

NOTE. According to LEGON *et al.* (2009) *S. fuscum* and *S. bovista* are synonyms; the species is mentioned as a separate one by WOJEWODA (2003) and red-listed in Poland (WOJEWODA & ŁAWRYNOWICZ 2006).

*Suillus bovinus* (L.) Roussel – on soil, under pine; J: 13, 1, 6, phases 6-6/7, X 1995, XI 1996.

*Suillus luteus* (L.) Roussel – on soil, under pine; J: 67, 84, 61, 85, 7, 13, 1, 6, 11, phases 4-8, IX-X 1994, VIII-XI 1995, VII-XI 1996, VII 1998.

*Suillus variegatus* (Sw.) Kuntze – on soil, under pine; CzO: 7, phase 9/10, X 1996.

*Tylopilus felleus* (Bull.) P. Karst. – on soil; CzO: 5, 6, 8, phase 9/10, VIII 1995, VII 1997.

*Xerocomus badius* (Fr.) E.-J. Gilbert [= *Boletus badius* (Fr.) Fr.] – on soil, under pine; J: 7, 1, 6, phases 6-6/7, IX-X 1995, VIII, X 1996, VII 1998; CzO: 2, 5, 6, 7, 8, phase 9/10, IX-X 1994, X-XI 1995, X-XI 1996, VII 1997.

*Xerocomus chrysenteron* (Bull.) Quél. [= *Boletus chrysenteron* Bull.] – on soil, under birch, pine; J: 85, phase 5, X 1995.

*Xerocomus subtomentosus* (L.) Quél. [= *Boletus subtomentosus* (L.) Quél.] – on soil, under birch, pine; J: 67, 84, 61, 85, phase 4-5, IX 1994, VII 1997, VII 1998.

### Cantharellales

*Botryobasidium vagum* (Berk. & M.A. Curtis) D.P. Rogers – on fallen branches of pine; CzO: 2, 5, 7, phase 9/10, IX 1994, VI, IX-X 1995, VI 1996; R; *det. W. Wojewoda*.

*Cantharellus cibarius* Fr. – on soil, under pine; J: 16, phase 8, VI, IX-X 1994, VI, VIII-X 1995, VI-VIII, X-XI 1996; CzO: 2, 5, 6, 7, phase 9/10, X 1994, VI, IX-X 1995, VIII, X-XI 1996, VII 1997, VII 1998; I.

*Tulasnella pallida* Bres. – on stump of pine; J: 11, phase 8, X 1995; *det. W. Wojewoda*.

### Gaeastrales

*Gaeastrum quadrifidum* Pers. – on soil; J: 11, 16, phase 8, VI, IX 1994; R.

*Sphaerobolus stellatus* Tode – on fallen twigs, on pieces of bark; J: 84, 7, 11, 16, phases 4, 6, 8, IX-X 1994, IX-X 1995.

### Gomphales

*Ramaria cf. abietina* (Pers.) Quél. – on soil; J: 16, phase 8, IX 1994, IX 1995.

*Ramaria eumorpha* (P. Karst.) Corner – on soil; J: 67, 7, 13, 6, phases 4, 6-6/7, IX-X 1994, IX 1995, VIII, X 1996.

### Hymenochaetales

*Coltricia perennis* (L.) Murril – on soil; J: 84, 61, 16, phases 4, 5, 8, IX 1994, X 1995, VI, X 1996, IX 1997; CzO: on forest path, phase 9/10, VII 1998.

*Hymenochaete tabacina* (Sowerby) Lév. – on fallen twigs of aspen; J: 11, 16, phase 8, IX 1994, VI, VIII 1995, VI, XI 1996; R; *det. E. Parmasto*.

*Hyphodontia aspera* (Fr.) J. Erikss. – on log and fallen branches of pine; J: 11, phase 8, VIII-IX 1995, CzO: 2, 8, phase 9/10, IX 1994, VIII, X 1995; *det. W. Wojewoda*.

*Hyphodontia quercina* (Pers.) J. Erikss. – on fallen branch of aspen; J: 16, phase 8, VI 1995; *det. W. Wojewoda*.

*Hyphodontia subalutacea* (P. Karst.) J. Erikss. – on fallen branches of pine; CzO: 6, 8, phase 9/10, IX 1994, X 1995, VI 1996.

*Inonotus rheades* (Pers.) Bondartsev & Singer – on dead trunks of aspen; J: 11, 16, phase 8, VI-X 1994, VIII, X 1995, VI-VII, X-XI 1996; CzO: 5, phase 9/10, V 1996; V; *rev. W. Wojewoda*.

*Phellinus punctatus* (Fr.) Pilát – on dead trunks of aspen; J: 11, 16, phase 8, X 1994, XI 1996; *rev. W. Wojewoda*.

*Rickenella fibula* (Bull.) Raithelh. – on soil, among mosses; J: 67, 84, 61, 85, 7, 13, 1, 6, phases 4-6/7, IX-X 1994, IX-XI 1995, VII-XI 1996.

*Schizopora paradoxa* (Schrad.) Donk [= *Hyphodontia paradoxa* (Schrad.) E. Langer & Vesterholt s.l.] – on dead trunk of aspen; J: 16, phase 8, X 1994.

*Tubulicrinis subulatus* (Bourdote & Galzin) Donk – on decayed woody debris of aspen; J: 11, phase 8, VI 1995; CzO: 2, phase 9/10, V 1996; *det. W. Wojewoda*.

### Polyporales

*Byssomerulius corium* (Pers.) Parmasto – on decayed woody debris; J: 11, 16, phase 8, IX-X 1994, XI 1997; *det. E. Parmasto*.

*Daedaleopsis confragosa* (Bolton) J. Schröt. – on dead trunks and branches of aspen; J: 11, 16, phase 8, VIII-X 1994, X 1995, VIII-XI 1996.

*Fomes fomentarius* (L.) J.J. Kickx – on dead trunk of birch; J: 16, phase 8, 1994, 1995, 1996, IX 1997.

*Gloeoporus taxicola* (Pers.) Gilb. & Ryvarden – on dead branches of pine; J: 11, phase 8, VIII 1995, CzO: 2, 7, phase 9/10, VIII 1994, VI 1995; R; *rev. W. Wojewoda*.

*Hyphoderma setigerum* (Fr.) Donk – on fallen twigs of aspen; J: 11, 16, phase 8, IX-X 1994, VI 1995; CzO: 8, phase 9/10, IX 1994; *det. W. Wojewoda*.

*Oligoporus alni* (Niemelä & Vampola) M. Piątek [= *Postia subcaesia* ss. Polish authors] – on log; J: 11, phase 8, IX-X 1994, X 1995, X 1996. See notes in WOJEWODA (2003; p. 474)).

*Panus suavissimus* (Fr.) Singer [= *Lentinus suavissimus* Fr.] – on fallen twigs of aspen; J: 16, phase 8, VI, IX 1994, IX 1995.

*Phanerochaete sanguinea* (Fr.) Pouzar – on fallen branches and logs of pine; J: 7, 1, 6, phases 6-6/7, X 1994, VI, X 1995; CzO: 2, 5, 8, phase 9/10, VIII 1994, VI, VIII, X 1995; *det. W. Wojewoda*.

*Phlebia rufa* (Pers.) M.P. Christ. – on stump of birch; J: 11, 16, phase 8, IX-X 1994, VI, VIII 1995, XI 1996; R; *rev. W. Wojewoda*.

*Phlebiella pseudotsugae* (Burt) K.H. Larss. & Hjortstam – on rotting log of pine; CzO: 2, phase 9/10, V 1995; *det. W. Wojewoda*.



*Phlebiella sulphurea* (Pers.) Ginns & M.N.L. Lefebvre [= *Phlebiella vaga* (Fr.) P. Karst.] – on fallen branches, on pieces of bark; J: 11, phase 8, IX-X 1995; CzO: 2, 5, 6, 7, 8, phase 9/10, IX 1994, VI, VIII, IX-X 1995, VI 1996, IX 1997; det. E. Parmasto.

*Polyporus ciliatus* Fr. – on fallen branches; J: 85, 6, phases 5, 6/7, VI 1994, VI 1996.

*Steccherinum fimbriatum* (Pers.) J. Erikss. – on rotting wood; J: 16, phase 8, X 1994, VI, VIII 1995, V 1996; R.

*Trametes ochracea* (Pers.) Gilb. & Ryvarden [= *Trametes zonatella* Ryvarden] – on fallen branches; J: 16, phase 8, IX-X 1994.

*Trichaptum fuscoviolaceum* (Ehrenb.) Ryvarden – on log of pine; J: 7, phase 6, VI, IX-X 1994.

### Russulales

*Amylostereum laevigatum* (Fr.) Boidin – on dead branches of *Juniperus*; J: 1, 11, 16, phases 6/7-8, X 1994, VI 1995; det. E. Parmasto.

*Auriscalpium vulgare* Gray – on buried cones of pine; J: 1, 6, 11, phases 6/7, 8, IX-X 1994, IX-X 1995, VII, IX, XI 1996, IX 1997; CzO: 2, 5, 6, 8, phase 9/10, VIII-IX 1994, IX, XI 1995.

*Gloeocystidiellum porosum* (Berk. & M.A. Curt.) Donk – on fallen branches of aspen; J: 11, phase 8, IX-X 1994, VIII 1995; det. W. Wojewoda.

*Lactarius helvus* (Fr.) Fr. – on soil; J: 1, phase 6/7, IX 1997.

*Lactarius hyssiginus* (Fr.) Fr. – on soil; J: 6, phase 6/7, XI 1997; V.

*Lactarius necator* (Bull.) Pers. [= *Lactarius turpis* (Weinm.) Fr.] – on soil, under birch; J: 11, phase 8, VIII 1996.

*Lactarius quieticolor* Romagn. – on soil, under pine; J: 61, phase 5, X-XI 1995, VIII 1996, VII 1998.

*Lactarius rufus* (Scop.) Fr. – on soil, under pine; J: 7, 13, 1, 6, 11, phases 6-8, IX-X 1994, IX-XI 1995, VII-VIII, IX-X 1996; CzO: 2, 5, 7, 8, phase 9/10, IX 1994, X 1995, VIII 1996, IX 1997.

*Lactarius tabidus* Fr. [= *Lactarius thejogalus* (Bull.) Gray s. auct.] – on soil; CzO: 5, phase 9/10, IX 1994, X 1995.

*Lactarius torminosus* (Schaeff.) Pers. – on soil, under birch; J: 11, 16, phase 8, IX-X 1994, X 1995, VII 1998.

*Lactarius uvidus* (Fr.) Fr. – on soil, under aspen; J: 85, 11, 16, phases 5, 8, X 1995.

*Lactarius vietus* (Fr.) Fr. – on soil, under birch; J: 84, phase 4, X 1995.

*Peniophora cinerea* (Pers.) Cooke – on fallen twigs of aspen; J: 11, phase 8, VI 1995.

*Peniophora polygonia* (Pers.) Bourdot & Galzin – on fallen twigs of aspen; J: 11, 16, phase 8, VI, IX-X 1994, V-VI, X 1995, XI 1996; det. W. Wojewoda.

*Russula amethystina* Quél. – on soil, under pine; CzO: phase 9/10, XI 1997, VII, IX 1998.

*Russula badia* Quél. – on soil, under pine; CzO: 2, 5, 6, 7, 8, phase 9/10, IX-X 1994, IX-XI 1995, X 1996.

*Russula clavipes* Velen. [= *Russula xerampelina* (Schaeff.) Fr. var. *elaeodes* Bres., *R. elaeodes* (Bres.) Bon.] – on soil under birch, aspen J: 84, 85, 7, 11, 16, phases 4-6, 8, X 1994, VIII, X 1995, VII-VIII 1996, VII 1997, VII 1998; CzO: 5, phase 9/10, IX 1997.

*Russula decolorans* (Fr.) Fr. – on soil, under pine; CzO: 2, 5, 6, 7, 8, phase 9/10, VIII, X 1995, VII-X 1996, VII 1997, VII 1998.

*Russula emetica* (Schaeff.) Pers. – on soil, under pine; J: 84, phase 4, X 1995; CzO: 2, 5, 6, 7, IX-X 1995, X 1996.

*Russula fragilis* (Pers.) Fr. – on soil, under aspen, pine, among mosses; J: 67, 84, 85, 7, 11, 16, phases 4-6, 8, IX-X 1994, VIII-X 1995, VII-VIII, X 1996, VII 1997, VII 1998; CzO: 2, 5, phase 9/10, X 1995, X 1996.

*Russula galochroa* (Fr.) Fr. [= *R. anatina* Romagn.] – on soil, under birch, aspen; J: 84, 85, 11, 16, phases 4-5, 8, IX 1994, VIII, X 1995, VII-VIII 1996, VII 1997, VII 1998.

*Russula* cf. *mustelina* Fr. – on soil, under pine; J: 16, phase 8, IX-X 1994, VII 1996.

*Russula nigricans* (Bull.) Fr. – on soil; CzO: phase 9/10, IX 1998.

*Russula paludosa* Britzelm. – on soil, under pine; J: 16, phase 8, IX-X 1994, VIII 1995, VII 1997; CzO: 2, 5, 6, 7, 8, IX 1994, VI, VIII-X 1995, VII-VIII 1996, VII, IX 1997, VII 1998.

*Russula subfoetens* Wm.G. Sm. – on soil, under birch; J: 84, 85, phases 4-5, VIII, X 1995.

*Russula turci* Bres. s. Maire – on soil, under pine; CzO: 7, phase 9/10, X 1995.

*Russula versicolor* Jul. Schäff. – on soil, under birch; J: 16, phase 8, IX 1994.

*Russula vesca* Fr. – on soil; CzO: 5, 8, phase 9/10, X 1995, IX 1997.

*Russula vinosa* Lindblad – on soil, under pine, among mosses; CzO: 5, phase 9/10, VI, X 1995, VII 1996, VII 1997.

*Russula* cf. *vinosobrunnea* (Bres.) Romagn. – on soil; J: 85, phase 5, X 1995.

*Russula xerampelina* (Schaeff.) Fr. s.str. – on soil, under pine; CzO: 5, 7, 8, phase 9/10, X 1995.

*Stereum hirsutum* (Willd.) Gray – on fallen branches of birch; J: 11, 16, phase 8, IX-X 1994, IX-X 1995, X-XI 1996.

*Vesiculomyces citrinus* (Pers.) E. Hagstr. [= *Gloiothele citrina* (Pers.) Ginns & G.W. Freeman] – on woody debris; J: 11, phase 8, IX 1994, X 1995; *det. W. Wojewoda*.

### **Thelephorales**

*Sarcodon squamosus* (Schaeff.) Quél. – on soil, under pine; J: 13, 1, 6, phases 6-6/7, IX-X 1994, X 1995, X-XI 1996.

*Thelephora caryophyllea* (Schaeff.) Pers. – on soil; J: 84, 1, 11, 16, phases 4, 6/7, 8, IX-X 1994, IX, XI 1995, VII 1997, VII, IX 1998; V.

*Thelephora palmata* (Scop.) Fr. – on soil (forest path); CzO: phase 9/10, VII 1997.

*Thelephora terrestris* Ehrh. – on soil, on woody debris; J: 67, 84, 61, 85, 7, 13, 1, 6, 11, phases 4-8, VI, IX-X 1994, VI, VIII-XI 1995, VI, VIII-XI 1996; CzO: 2, 5, 6, 7, 8, phase 9/10, IX 1994, IX 1995, IX 1996.

### **Agaricomycetes, incertae sedis**

*Peniophorella praetermissa* (P. Karst.) K.-H. Larss. [= *Hyphoderma praetermissum* (P. Karst.) J. Erikss. & Å. Strid] – on fallen branches of pine, aspen; J: 11, phase 8, IX 1994, VI 1995; *det. W. Wojewoda*.

*Resinicium bicolor* (Alb. & Schwein.) Parmasto – on woody debris; J: 16, phase 8, VI 1995; *det. W. Wojewoda*.



## THE POLISH BOTANICAL SOCIETY

announces the

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*„Planta in vivo, in vitro et in silico”*

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Warsaw University of Life Sciences-SGGW

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55th PBS Meeting Office

Katarzyna Marciszewska, Ph.D.

**Forest Botany Division, Forestry Faculty Warsaw University of Life Sciences-SGGW**

159 Nowoursynowska St., 02-776 Warsaw, Poland

E-mail: [ptb.marciszewska@wl.sggw.pl](mailto:ptb.marciszewska@wl.sggw.pl)

Tel. (+4822) 5938026 Fax. (+4822) 5938022

<http://www.zjazd55.ptb.waw.pl>