



DOI: <https://doi.org/10.23859/estr-240405>

EDN: <https://elibrary.ru/opxmx>

UDC 631.466.1*630*114

Article

Fungi of forest litter in experimental plantations of lodgepole pine (*Pinus contorta* Dougl.)

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Abstract. The paper deals with the community of soil fungi of forest litter in experimental plantations of the Krasnozatskiy forestry in the Komi Republic, created in a clear-cut area using an introduced species, i.e. lodgepole pine (*Pinus contorta* Dougl.), during the growing season of 2023. The species composition of fungi is shown to include 34 species from 8 genera, 7 orders and 2 divisions, including sterile mycelium. Most species belong to ascomycetes (88.2% of the total species diversity). Representatives of the genus *Penicillium* dominate by species number, frequency of occurrence and abundance. The litter fungal community is subject to significant seasonal dynamics with the microbial abundance increasing from May ($17.1 \pm 7.5 \times 10^3$ CFU/g) to September ($60.9 \pm 19.2 \times 10^3$ CFU/g) and the number of species increasing from 12 to 25. The biomass of fungi during the growing season ranges from 1.04 ± 0.17 to 7.68 ± 3.31 mg/g. The results obtained can be further used in assessing the impact of forest crops grown with the use of introduced species on the forming forest litter.

Keywords: pine forests, experimental plantations, introduced species, fungi, microbial biomass

Funding. The work was carried out with the financial support of the state tasks of the Institute of Biology, Komi Science Centre of the Ural Branch of the Russian Academy of Sciences titled “Environment-forming role and productivity of forest and peatland ecosystems of the European North-East of Russia” (№ 125020501547–8) and “Microbial communities of ecosystems in the North, their biotechnological potential and its realisation technologies” (№ 125021201993–3).

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To cite this article: Kovaleva, V.A. et al., 2025. Fungi of forest litter in experimental plantations of lodgepole pine (*Pinus contorta* Dougl.). *Ecosystem Transformation* **8** (3), 82–98. <https://doi.org/10.23859/estr-240405>

Received: 05.04.2024

Accepted: 02.08.2024

Published online: 01.08.2025

DOI: <https://doi.org/10.23859/estr-240405>EDN: <https://elibrary.ru/opxmxh>

УДК 631.466.1*630*114

Научная статья**Микромицеты лесной подстилки
в опытных культурах сосны скрученной
(*Pinus contorta* Dougl.)**В.А. Ковалева , Ю.А. Виноградова , Т.А. Пристова* *Институт биологии Коми научного центра Уральского отделения РАН, 167000, Россия,
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Аннотация. В течение вегетационного периода 2023 г. изучено сообщество почвенных микромицетов лесной подстилки в экспериментальных культурах Краснозатонского лесничества Республики Коми, созданных на вырубке с применением вида-интродуцента – сосны скрученной (*Pinus contorta* Dougl.). Показано, что видовой состав микромицетов насчитывает 34 вида из 8 родов, 7 порядков и 2 отделов, включая стерильный мицелий. Большинство видов относится к аскомицетам (88.2% от всего видового разнообразия). Представители рода *Penicillium* доминируют по количеству видов, частоте встречаемости и обилию. Микоценоз подстилки подвержен значительной сезонной динамике, численность микромицетов возрастает с мая ($17.1 \pm 7.5 \times 10^3$ КОЕ/г) по сентябрь ($60.9 \pm 19.2 \times 10^3$ КОЕ/г), количество видов увеличивается с 12 до 25. Биомасса микромицетов за вегетационный период варьирует от 1.04 ± 0.17 до 7.68 ± 3.31 мг/г. Полученные результаты в дальнейшем могут быть использованы при оценке влияния лесных культур, созданных с применением интродуцентов, на формирующуюся лесную подстилку.

Ключевые слова: сосновые леса, экспериментальные культуры, интродуцент, микоценоз, микробная биомасса

Финансирование. Работа выполнена при финансовой поддержке тем госзадания Института биологии Коми НЦ УрО РАН «Средообразующая роль и продуктивность лесных и болотных экосистем европейского северо-востока России», 2025–2029 гг. (№ 125020501547-8) и «Микробные сообщества экосистем Севера, их биотехнологический потенциал и технологии его реализации» регистрационный номер (№ 125021201993-3).

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Для цитирования: Ковалева, В.А. и др., 2025. Микробиоты лесной подстилки в опытных культурах сосны скрученной (*Pinus contorta* Dougl.). *Трансформация экосистем* 8 (3), 82–98. <https://doi.org/10.23859/estr-240405>

Поступила в редакцию: 05.04.2024

Принята к печати: 02.08.2024

Опубликована онлайн: 01.08.2025

Introduction

Intensive use and reproduction of forests requires an accelerated timber growing that implies, among other things, the use of fast-growing tree species. One of such species for the taiga zone of the Russian Federation is lodgepole pine (*Pinus contorta* Dougl.) (Melekhov, 1984). The natural range of this species is the western part of North America (Elias, 2014). Lodgepole pine is known to be superior to Scots pine by the growth rate (Fedorkov and Gutiy, 2017; Feklistov et al., 2008; Elfving et al., 2001; Raevsky, 2015). The use of introduced species can lead not only to their naturalisation and introduction into natural communities (Zagurskaya, 2022), but also to changes in the quality of leaf waste and litter during artificial forest restoration. This, in turn, can cause changes in ecosystem processes (decomposition and nutrient cycling rates), as well as in ecosystem properties (changes in soil carbon pool values) (McIntosh et al., 2012). Even if introduced and native species are functionally similar but differ by their growth rate and, consequently, by input of organic material from leaf fall into the litter, this will still result in changes in ecosystem properties and processes (McIntosh et al., 2012).

The process of forest ecosystem restoration affects the intensity of biological processes and organic matter accumulation. Soil microorganisms determine the direction and rate of plant waste transformation (Dobrovolskaya et al., 2015). Soil fungi possess a wide range of extracellular enzymes, produce acids, antibiotics, toxins, phytohormones, pigments, etc. (Berestettsky, 2008; Dobrovolsky and Nikitin, 2006; Karlovsky, 2008; Looby and Treseder, 2018). Due to this, they are actively involved in the processes of humus formation, immobilisation and mobilisation of nutrients and so maintain the homeostasis and recovery under anthropogenic disturbances in forest ecosystems (Aleksandrova et al., 2006a; Bakhmet and Medvedeva, 2022; Dobrovolskaya et al., 2015; Habibullina et al., 2018). Forest litter has optimal conditions for the development of different groups of fungi: high organic matter content, favourable physico-chemical and hydrothermal parameters (Dobrovolskaya et al., 2015; Kitikidou, 2012). The fungal community of forest litter includes fungal species of phyllosphere and just soil fungus, which determines high diversity and changes in taxonomic structure of fungi during the decomposition of plant waste (Habibullina and Tvorozhnikova, 2007; Kurakov and Semenova, 2016; Pristova et al., 2012; Semenova, 2002; Terekhova et al., 1998; Voříšková and Baldrian, 2013). The fungal communities of forest plantations differ from those of natural stands qualitatively and quantitatively, which roots in the characteristic features of forest litter between these communities (Shebalova, 2008; Shebalova and Zalesov, 2006). Litter fungi in pine plantations, as well as in forest plantations created using introduced species, are rarely studied (Bakhmet and Medvedeva, 2022; McIntosh et al., 2012; Shebalova and Zalesov, 2006). We did not find the study results on soil fungal community in lodgepole pine plantations in the literature.

The aim of the research was to assess the species diversity of fungi, their number and biomass in experimental plantations of lodgepole pine.

Materials and methods

The study materials were experimental plantations of lodgepole pine located on the territory of the Krasnozatonksy district forestry (Division 34) of the Syktyvkar forestry in the Komi Republic (N 61°40' E 51°03') (Fig. 1). Forest litter fungi were studied during the growing season of 2023.

Forest plantations were planted as two-year-old bare-root seedlings, grown from seeds of Swedish reproduction, instead of logged cowberry-lichen pine forest in 2006. The soil was prepared by the tractor-mounted plow TK-1 in summer 2005. The site area is 1.0 ha, total number of seedlings – 2531 units. The soil type is illuvial-iron podzol. The average litter thickness is 3.9 cm. The site hosts 36 plant species: 8 species of trees, 7 species of shrubs and dwarf shrubs, 16 species of herbs, 5 species of mosses, and 3 species of lichens. Tree and shrub species include *Betula pendula* Roth., *B. pubescens* Ehrh. and single specimens of *Salix caprea* L., *Sorbus aucuparia* L., *Populus tremula* L., and *Picea obovata* Ledeb. of seed origin.

The total projective cover (TPC) of ground cover plants at the site averages 90%, varying from 60 to 100%. The projective cover of moss cover is 55% dominated by *Polytrichum commune* Hedw.; *Pleurozium schreberi* (Willd ex Brid.) Mitt.; *Dicranum polysetum* Sw. The dominating position of mosses, in particular polytrichum mosses, is related to the microrelief specificity: a combination of rows with pine plantings and inter-rows (depressions). Moist inter-rows favours the development of moss cover. The projective cover of the herbaceous-dwarf shrub layer averages 45%, varying from 15 to 80%. Dwarf shrubs play a leading role in formation of the herbaceous-dwarf shrub layer, their projective cover can reach 55% of TPC, averaging 30% with dominating *Vaccinium vitis-idaea* L. Among herbs, *Calamagrostis epigeios* (L.) Roth.; *Chamerion angustifolium* (L.) Scop.; *Luzula pilosa* (L.) Willd; *Agrostis gigantea* Roth. are common species. In the living soil ground cover, ferns, lycopods, and lichens are rare or single with projective cover of less than 1%.



Fig. 1. Experimental crops of lodgepole pine in the Krasnozatonksy forestry.

The litter fungal community under lodgepole plantations was analysed in the beginning (May) and the end (August and September) of the vegetation period. Forest litter samples were collected from the entire area of experimental cultures: 12 samples in May, 12 samples in August, and 15 samples in September. The material was sampled according to the generally accepted methods (Kurakov, 2001). Prior to microbiological analyses, collected litter samples were stored in a freezer at $-18...-20$ °C. Isolation of fungi and their number calculation (in colony-forming units per gram of absolutely dry soil – CFU/g a.d.s.) were done by sowing from serial dilutions onto the agar Chapek's medium (pH = 4.5) in fivefold repetition for each sample. Calculation of fungal colonies was carried out 5–7 days later (Kazeev et al., 2003). Taxonomic affiliation of fungal isolates was identified to species using identification guides of fungi (Alexandrova et al., 2006b; Domsch et al., 2007; Egorova, 1986; Ellis, 1971; Pitt, 1991; Ramirez, 1982, etc.). Name and systematic position of taxa were unified using the database – Index Fungorum¹. To characterise the complex of cultivated soil fungi, we applied the Shannon species diversity index (H), the Pielou index of evenness (E), the Simpson domination index (D), as well as the spatial frequency and the relative abundance indices of species (Kurakov, 2001). Spatial frequency was estimated as the proportion of samples, where the particular species was found, to the total number of collected samples.

Spore number, length and viability of fungal mycelium, the fungal biomass value was estimated by the luminescence microscopy (Metody..., 1991; Polyanskaya, 1996) using fluorescein diacetate (FDA) dye (Gaspar et al., 2001).

Statistical processing of the obtained data was carried out using Microsoft Excel 2010 programme.

Results and discussion

Number and species diversity of fungi

The number of fungi in the studied samples differed significantly in May and September. The fungal number was $17.1 \pm 7.5 \times 10^3$ CFU/g in May but $25.4 \pm 14.3 \times 10^3$ and $60.9 \pm 19.2 \times 10^3$ CFU/g in August and September, respectively. The number of estimated fungi is a relatively low value as compared to natural pine forests on the Kola Peninsula where the number of fungi varies from 46×10^3 to 216×10^3 CFU/g (Korneikova et al., 2018).

The litter samples of the studied pine plantation were isolated for 34 species of fungi from 8 genera, 7 orders and 2 divisions, including sterile mycelium (Table 1).

By the taxonomic structure analysis of the fungal community, the species diversity of fungi in litter under experimental lodgepole pine plantation exceeds the results we obtained earlier for the lodgepole pine plantations in the Storozhevsk forestry – 17 species from 9 genera (Kovaleva et al., 2023) and for the pine forests on the Kola Peninsula – 12 species from 5 orders (Korneikova et al., 2018).

The Mucoromycota division is not numerous and includes three species as *Mortierella alpina*, *Mucor hiemalis*, and *Umbelopsis vinacea* that make 8.8% of the total species composition. These species are widely distributed and are common soil representatives not only in the middle taiga of the Komi Republic (Khabibullina and Tvorozhnikova, 2007; Khabibullina et al., 2018), but also in higher latitudes (Vinogradova et al., 2019, 2023). *M. hiemalis*, like other representatives of the *Mucoromycota* division, belongs to the group of saprotrophic fungi that first utilise readily-available sugars (Domsh et al., 2007). For this reason, it is normally found in litter towards the end of the growing season at the fresh leaf fall stage (Table 1). The species prefers plant waste from the herb-dwarf shrub layer.

The main part of fungal species belong to anamorphic ascomycetes (from the Ascomycota division) – 30 species from 5 genera that make 88.2% of the total species composition. Fungi of the *Penicillium* genus, which is dominant in soils of the Komi Republic (Khabibullina et al., 2018; Kovaleva et al., 2023; Sizonenko et al., 2010; Vinogradova et al., 2019, 2023), account for 51% of the total number of isolated species (Fig. 2).

Fungi of the *Penicillium* genus belong mainly to typical hydrolytics that can develop on poor-available nutrients and assimilate difficult-to-decompose polymeric substrates at places with low rate mineralisation processes (Domssh et al., 2007; Osono, 2007; Vinogradova et al., 2023). In this connection, the litter micromycete complex in the early vegetation period is mostly composed of fungi of the *Penicillium* genus that participate in mineralisation of the previous year's fall.

¹ Index Fungorum. Web page. URL: <https://www.indexfungarum.org> (accessed: 14.03.2024).

Table 1. Species diversity, relative abundance of species and spatial frequency of occurrence (SFO) of soil fungi species. Frequency of occurrence categories: > 60% dominant (D), 30–60% – often-met (OM), 10–30% – rare (R), < 10% – occasional (O).

Species	May		August		September	
	Abundance, %	SFO	Abundance, %	SFO	Abundance, %	SFO
Division Mucoromycota						
Order Mortierellales						
<i>Mortierella alpina</i> Peyron	–	–	1.1	O	–	–
Order Mucorales						
<i>Mucor hiemalis</i> Wehmer	–	–	0.5	O	2.5	R
Order Umbelopsidales						
<i>Umbelopsis vinacea</i> (Dixon.) W. Gams	–	–	0.5	O	0.3	O
Division Ascomycota						
Order Capnodiales						
<i>Cladosporium herbarum</i> (Pers.) Link	–	–	–	–	0.5	O
Order Eurotiales						
<i>Penicillium aurantiogriseum</i> Dierckx	–	–	5.3	R	0.3	O
<i>P. brevicompactum</i> Dierckx	–	–	–	–	0.3	O
<i>P. canescens</i> Sopp	–	–	5.3	OM	15.4	OM
<i>P. camemberti</i> Thom	–	–	–	–	0.3	O
<i>P. commune</i> Thom	–	–	–	–	0.5	R
<i>P. citreonigrum</i> Dierckx	–	–	0.5	O	2.2	OM
<i>P. expansum</i> Link	1.6	O	2.1	O	–	–
<i>P. decumbens</i> Thom	1.6	O	4.3	OM	8.0	Д
<i>P. dierckxii</i> Biourge	–	–	1.1	O	1.1	R
<i>P. granulatum</i> Bainier	–	–	6.9	R	–	–
<i>P. glabrum</i> (Wehmer) Westling	–	–	0.5	O	–	–
<i>P. jensenii</i> K.W. Zaleski	6.3	R	0.5	O	6.3	OM
<i>P. lanosum</i> Westling	–	–	0.5	O	3.3	R
<i>P. lividum</i> Westling	3.2	R	2.7	R	5.5	D
<i>P. raistrickii</i> G. Sm.	–	–	0.5	O	1.4	R
<i>P. simplicissimum</i> (Oudem.) Thom	–	–	2.1	O	2.7	R

Species	May		August		September	
	Abundance, %	SFO	Abundance, %	SFO	Abundance, %	SFO
<i>P. thomii</i> Maire	15.9	OM	36.2	D	28.6	D
<i>P. vulpinum</i> (Cooke & Masee) Seifert & Samso	1.6	O	6.9	R	–	–
<i>P. waksmanii</i> K.M. Zaleski	1.6	O	–	–	–	–
<i>Penicillium</i> sp.	3.2	R	1.1	R	1.9	R
<i>Talaromyces diversus</i> (Raper & Fennell) Samson, N. Yilmaz & Frisvad	–	–	0.5	O	-	-
<i>T. funiculosus</i> (Thom) Samson, N. Yilmaz, Frisvad & Seifert	–	–	–	–	3.0	OM
<i>T. rugulosus</i> (Thom) Samson, N. Yilmaz, Frisvad & Seifert	6.3	R	1.6	R	–	–
<i>T. variabilis</i> (Sopp) Samson, N. Yilmaz, Frisvad & Seifert	–	–	0.5	O	0.3	O
Order Hypocreales						
<i>Trichoderma koningii</i> Oudemans	1.6	O	–	–	–	–
<i>T. hamatum</i> (Bonord.) Bainier	–	–	2.7	R	–	–
<i>T. polysporum</i> (Link) Rifai	–	–	–	–	1.4	O
<i>T. viride</i> Pers.	1.6	O	0.5	O	0.5	O
Order Thelebolales						
<i>Pseudogymnoascus pannorum</i> (Link) Minnis & D.L. Lindner	–	–	–	–	14.0	R
Sterile mycelium						
<i>Mycelia sterilia</i>	55.6	D	15.4	OM	0.3	O

The *Trichoderma* and *Talaromyces* genera are represented by 4 species. The other genera are represented by single species. Fungi of the *Talaromyces* genus are normally soil saprotrophs, widely distributed (Domssh et al., 2007; Kurakov and Semenova, 2016). Species of the *Trichoderma* genus occur in various substrates (Alexandrova et al., 2006b) and secrete cellulases, which make them part of the group of cellulose-destroying fungi involved in the mineralisation of difficult-to-decompose organic compounds (Rakhleeva et al., 2011; Osono and Takeda, 2001; Osono et al., 2003). Species of the *Trichoderma* genus were isolated from litter samples at any sampling dates: in May – *T. koningii*, in August – *T. hamatum*, in September – *T. polysporum* and at any year time – *T. viride*.

Penicillium thomii dominates by the frequency of occurrence and abundance in litter samples at any sampling dates. The abundance of this species increases from May to September (Table 1). In addition to *P. thomii*, the species *Penicillium jensenii*, *P. lividum*, *Trichoderma viride* and *Mycelia sterilia* are common at any sampling dates. The relative abundance towards the end of the growing season increases for the *Penicillium decumbens* and *Penicillium lividum* species, whereas the abundance of the *Trichoderma viride* and *Mycelia sterilia* species decreases by September (Table 1). The latter actively

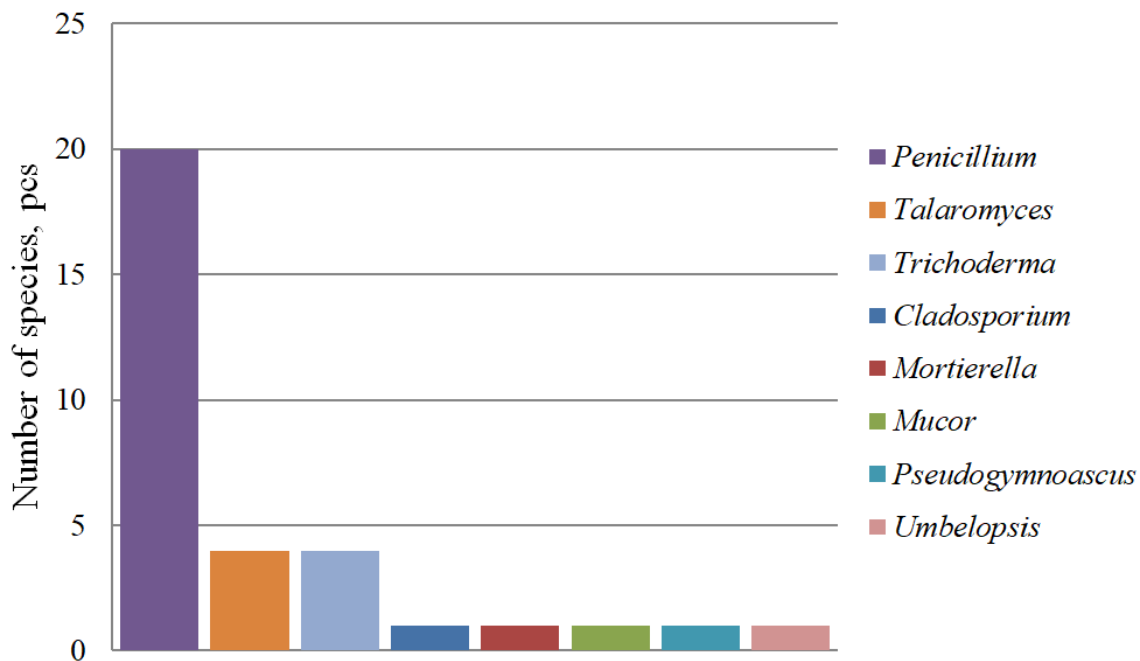


Fig. 2. Number of micromycete species of the main genera.

develop at the final stages of organic matter mineralisation and play an important role in decomposition of the previous year's fall (Rakhleeva et al., 2011; Osono, 2005; Osono et al., 2003). Sterile mycelium was isolated throughout the whole growing season with different frequency of occurrence and relative abundance values (Table 1). The formation of sterile mycelium results from the loss of the spore formation capacity on every nutrition medium used. The presence of various forms of sterile mycelium is a characteristic feature of soil micromycete communities in natural pine forests (Evdokimova and Mozgova, 2001; Korneikova et al., 2018).

Thus, the micromycete complex of the forest litter under lodgepole pine plantation consists of widely distributed species that inhabit both soils of the middle taiga of the Komi Republic (Khabibullina et al., 2018; Sizonenko et al., 2010) and far northward soils of higher latitudes (Vinogradova et al., 2019, 2023).

According to the seasonal observations of changes in the abundance and species diversity of soil fungi from lodgepole pine litter, the micromycete community had relatively low quantitative and qualitative indices in May. Twelve species of fungi from 3 genera, including sterile mycelium, were isolated from litter samples in May. The isolated species were ascomycetes, among them 8 species belonged to the *Penicillium* genus. *Penicillium thomii* and sterile mycelium dominated by the frequency of occurrence. Sterile mycelium demonstrated high values of relative abundance in May (Fig. 3).

The other species, according to the frequency of occurrence value, compose the group of rare and occasional species, the relative abundance of which varies from 1.6 to 6.3%. The processes of organic matter decomposition and transformation, in which fungi play a key role, are slowed down at the beginning of the vegetation period when the amount of fresh plant fall is low and the litter is not warm enough. The high abundance value of sterile mycelium also indicates unfavourable conditions for the development of litter fungi in May. Therefore, this period has a small group of fungi, represented by species of the *Penicillium* genus, participating in mineralisation of the last year's fall.

Towards the end of the vegetation period, the number of fungi and simultaneously the species diversity of mycocenosis increases. 24–25 species of fungi, including sterile mycelium, were isolated from the litter in August–September (Table 1). The group of dominant and frequently occurring species includes *Penicillium canescens*, *P. decumbens*, *P. thomii* and sterile mycelium in August. In September, the group of dominant and frequently occurring species increases significantly but mainly by the species of the *Penicillium* genus such as *Penicillium citreonigrum*, *P. jensenii*, *P. lividum* and additionally by *Talaromyces funiculosus*.

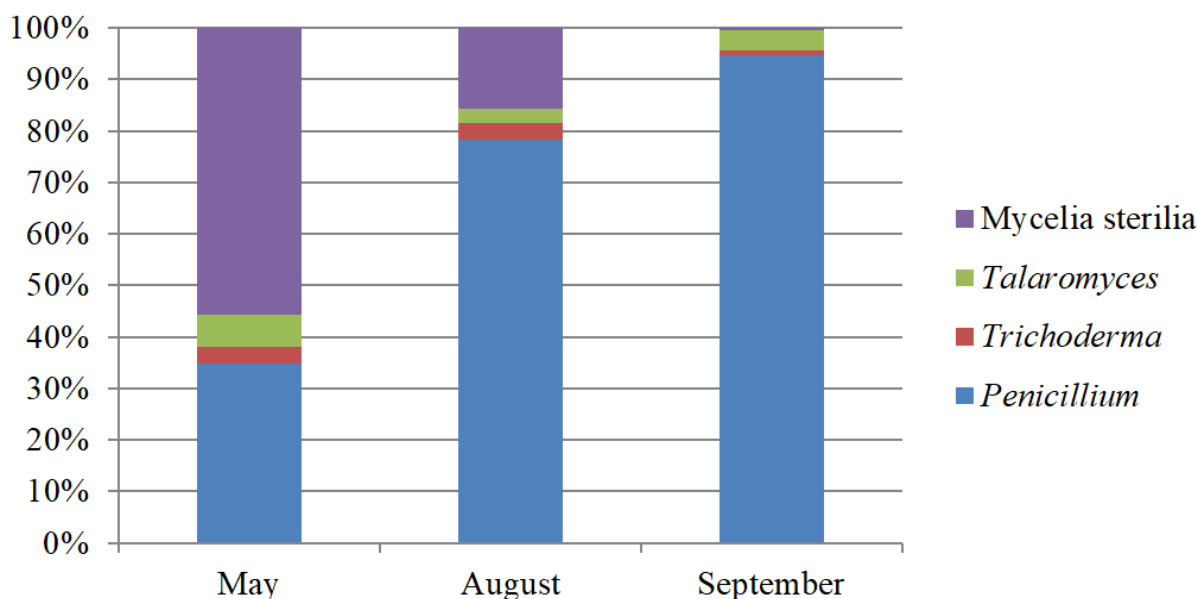


Fig. 3. Seasonal dynamics of relative abundance of sterile mycelium and the main fungal genera in litter.

In contrast to the early vegetation period, sterile mycelium disappears from the group of dominants at the end of the vegetation period. Its number and abundance sharply decrease (Fig. 3). At the same time, the majority of species also belong to the group of rare and occasional species (Table 1). Their development is associated with changes in the trophic structure of the micromycete community. The fresh plant fall stimulates the development of fungi and increases their species diversity by new species as *Penicillium aurantiogriseum*, *P. canescens*, *P. citreonigrum*, *P. dierckxii*, *P. lanosum*, *P. raistrickii*, *P. simplicissimum*, *Talaromyces funiculosus*, *T. variabilis*, *Trichoderma hamatum*, *T. polysporum*, *Umbelopsis vinacea* and others. Moreover, the number and species diversity of litter fungi in the studied artificial plantation increases at the late vegetation period both by soil species and plant species that attained the litter together with the fall (*Cladosporium herbarum*, *Mucor hiemalis*).

The end of the vegetation period is characterised by the maximum high values of the Shannon species diversity index ($H = 2.37$), which is associated with an increase in the number of dominant species and their relative abundance (Table 2).

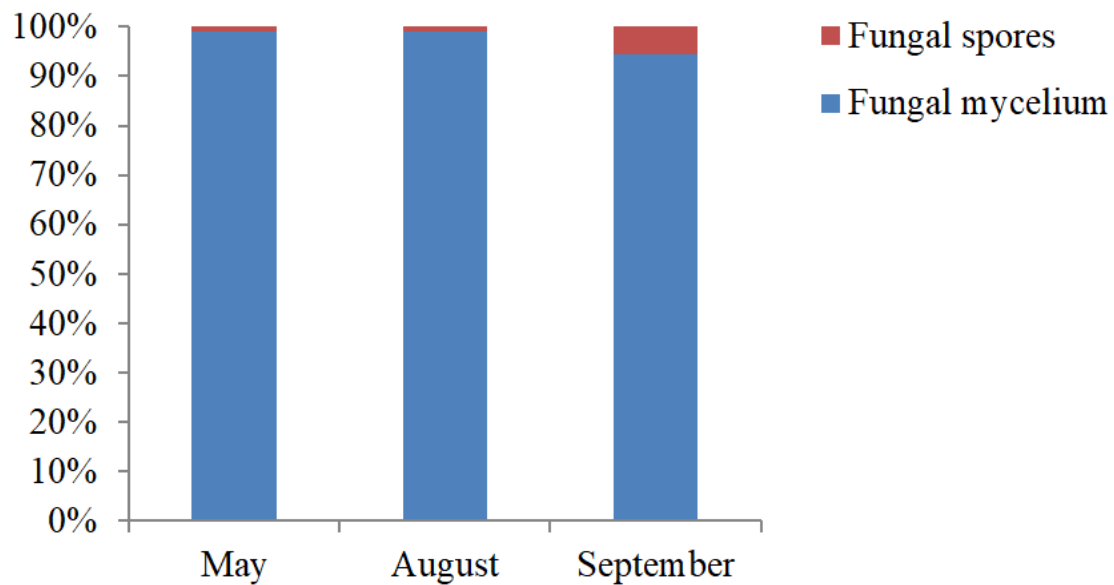
The obtained data on the relative abundance of species are also confirmed by the Simpson index. The Simpson index has its minimum value in September when the micromycete community is reorganised and increases in number of species dominating by relative abundance. At the same time, the evenness of the micromycete community increases.

Biomass of fungi

The fungal biomass in the litter of experimental plantation of lodgepole pine during the growing season decreases from to 7.68 ± 3.31 in May to 1.04 ± 0.17 mg/g in September. This is due to the fact that the main contribution to the biomass structure is made by fungal mycelium. Its length in May attains the maximum values, 1963 ± 865 m/g a.d.s., and decreases to 139.22 ± 21.56 m/g a.d.s. by the end of the growing season in September. The share of spores in fungal biomass structure increases in September, as compared to May (Fig. 4). In general, the observed biomass values of microscopic fungi are higher than our previously obtained data on fungal biomass in the litter of lodgepole pine from the Storozhevsk forestry (Kovaleva et al., 2023). In podzols of the Kola Peninsula, fungal biomass in natural pine forest ranges from 1.2 to 2.1 mg/g soil (Korneikova et al., 2018).

Table 2. Values of diversity, dominance and evenness indices for complexes of microscopic fungi.

Sampling month	Index		
	Shannon	Simpson	Pielou
May	1.58	0.33	0.73
August	2.32	0.17	0.78
September	2.37	0.14	0.81

**Fig. 4.** Ratio of fungal mycelium and spore biomass during the growing season.

Conclusion

The experimental site under study is a result of numerous transformations: from natural forest ecosystem (pine forest) and subsequent cutting to ploughing and planting of seedlings of native Scotch pine and introduced lodgepole pine. The destruction of forest litter followed by its restoration in the artificially created forest phytocenosis affects the mycocenosis. In general, the number of microscopic fungi in artificial forest is lower than that in natural pine forests. By the studied taxonomic structure of litter mycocenosis under lodgepole pine plantation, its species composition includes 34 species from 8 genera (including sterile mycelium) belonging to two divisions. The species diversity of fungi increases at the end of the growing season: 12 species from 3 genera of fungi were identified in May and 24 species from 7 genera – in September.

Most species belong to the *Penicillium* genus. Common for any sampling dates are *Penicillium thomii*, *P. jensenii*, *P. lividum*, *Trichoderma viride* and *Mycelia sterilia*. *Penicillium thomii* dominates by the frequency of occurrence and the relative abundance. Its abundance increases from May to September.

Viewing the seasonal dynamics, the fungal community changes its characteristics. In August–September, the fungal community changes its dominant groups with the increasing number of rare and occasional species, which may have been isolated from the litter in form of sterile mycelium at the beginning of the growing season. The end of the growing season is characterised by the development of saccharolytic fungi, which first inhabit fresh plant fall and decompose simple organic compounds. At the same time, biopolymer degraders and oligotrophic fungi undergo active development. They complete mineralisation of the last year's fall. Plant fungi serve an additional source of fungal diversity at the end

of the growing season. They enter the soil litter within plant fall. The number of fungi increases from $17.1 \pm 7.5 \times 10^3$ CFU/g in May to $60.9 \pm 19.2 \times 10^3$ CFU/g in September.

Fungi participating in the process of litter formation are characterised by comparatively high biomass indices. The maximum biomass development is observed at the beginning of the vegetation period (May) dominated by the fungal mycelium (by 99%). The high biomass of fungi at the beginning of the vegetation period against relatively low indices of species diversity and abundance is possibly associated with a decrease in intraspecific competition and abundant development of sterile mycelium.

According to the analysis of the obtained quantitative and qualitative indices and by the comparison with the data of other researchers (Evdokimova and Mozgova, 2001; Khabibullina and Tvorozhnikova, 2007; Khabibullina et al., 2018; Korneikova et al., 2018), the litter fungal community of lodgepole pine plantation is represented by the species typical of taiga natural forest ecosystems and cosmopolitan species occurring everywhere. This allows us to conclude that 20 years gone since forest plantation replaced once logged pine forest are enough for the forest litter to host the fungal community similar to natural forest ecosystems.

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