Ophiostomatoid fungi vectored by bark beetles and colonizing trees of *Pinus sylvestris* in Sumy region (Ukraine)

K. Davydenko¹, D. Baturkin²

Intensive mortality of *Pinus sylvestris* trees has recently been observed in the Sumy region in eastern Ukraine. There are two pine bark beetle species (*Ips acuminatus* and *Ips sexdentatus*), which spread resulted in considerable forest damage in Ukraine. The study of ophiostomatoid fungi vectored by bark beetles is very important to assess total harm of these insects.

Therefore, the aim of our research was i) to identify ophiostomatoid fungi associated with weakened and dying Scots pine trees infested by bark beetles in the Sumy region; ii) to test the pathogenicity of these ophiostomatoid fungi to evaluate their potential threat to Scots pine. The fungi were isolated from bark beetle galleries and identified based on morphological properties and DNA sequences.

In total, eight ophiostomatoid fungi (*Graphium sp.*, *Grosmannia sp.*1, *Ophiostoma bicolor*, *O. ips*, *O. canum*, *O. piceae*, *O. minus*, *Ophiostoma sp.*1) were isolated from Scots pine trees infested by bark beetles. Scots pine seedlings were inoculated with eight fungi and sterile medium (control) to evaluate their pathogenicity. The inoculated seedlings were examined finally in 6 month after inoculation. Inoculation with *O. minus* produced significantly largest lesions and only this fungus caused mortality of pine seedlings. In total, all eight fungal species inoculated caused resin exudation and staining the bark around inoculations in Scots pine seedlings and five fungi caused different rate of seedlings decline. The size of stained sapwood was also greater following *O. minus* inoculations than other fungi or the control. All ophiostomatoid fungi caused significantly longer necrotic lesions and more occlusions in the sapwood than the controls.

Therefore, based on the ability of various ophiostomatoid fungi to weaken and kill pine seedlings and stain sapwood, *O. minus* was the most dangerous species for Scots pine trees, followed by *Graphium sp.* and *Ophiostoma sp.*1. The occurrence of ophiostomatoid fungi in the sapwood of Scots pine is consistent with the concept of their primary role in the colonization of the fresh sapwood of trees in the succession of microorganisms during wood decay.

**Key words:** ophiostomatoid fungi; Scots pine; bark beetles; pathogenicity; forest health.

Introduction. Climate change worldwide has been linked to increased mortality of conifers due to drought, water stress, and consequently associated insect outbreaks, especially bark beetles (Cottrell et al., 2020; Dobor et al., 2020).

Some bark beetles can cause extensive harm to the spruce and pine forests across Europe including Ukraine. Thus, massive mortality of Scots pine (*Pinus sylvestris*) trees has been observed in the Sumy region in northern Ukraine since 2016 due to the drought and
subsequent bark beetle outbreak. There are now two
the most aggressive pine bark beetle, pine engraver
beetle *Ips acuminatus* (Gyllenhal, 1827; Curculionidae,
Scolytinae) and six-toothed bark beetle *Ips sexdentatus*
(Boerner, 1766; Curculionidae, Scolytinae), the spread of
which resulted in considerable damage of the
forest in Ukraine last decade (Meshkova et al., 2018; Meshkova & Borisenko, 2018). This complex
of bark beetles is capable of killing pine forests over
large areas in Ukraine and Europe (Meshkova et al.,
2018). Moreover, bark beetles carry microbial spores
on their exoskeleton, transferring them to trees during
the colonization or feeding in the phloem of pine trees
(Jankowiak, 2006; Bueno et al., 2010).

Therefore, now bark beetles are among the most
economically and ecologically important pests of
the world forests (Six, 2013; Cottrell et al., 2020;)
but greater problems arise from their associated
phytopathogenic blue-stain fungi, so-called
ophiostomatoid fungi. This group includes genera
such as *Ophiostoma*, *Ceratocystis*, *Graphilbum*,
*Raffaelea* and *Leptographium* in the *Ophiostomatales*,
and *Ceratocystis* and *Graphium* in the
*Microascales* (De Beer et al., 2013). The ophiostomatoid group
contains a lot of tree pathogens, which are responsible
for serious tree diseases, wood discoloration, and
even high rates of tree mortality (Davydenko et al.,
2017, 2019; Dori-Bachash et al., 2015; Repe et al.,
2015). Many studies demonstrated that in the case of
aggressive bark beetles that are able to kill healthy and
slightly weakened trees, phytopathogenic fungi may
help the beetles to overwhelm tree defenses (Six, 2003;
Six, 2013; Wingfield et al., 2017).

There is only a little information about the
association of ophiostomatoid fungi with bark beetles
and their pathogenicity to pine trees in Ukraine
(Davydenko et al., 2014; Davydenko et al., 2017).
In a recent study in Ukraine, six blue-stain fungi,
including, *O. cf. rectangulosporium*, *O. ips* *O. minus*,
*O. pallidulum*, *O. picea* and *O. olivaceum* were isolated
from the *Ips acuminatus* and their pathogenicity were
tested (Davydenko et al., 2017, 2019). Among these,
*O. minus* was the most pathogenic one when inoculated
into *P. sylvestris* trees that it was also demonstrated in
other studies (Jankowiak 2006; Jankowiak & Bilański,
2012).

Therefore, the aim of our research was 1) to identify
ophiostomatoid fungi associated with weakened and
dying Scots pine trees infested by bark beetles in the
Sumy region; 2) to test the pathogenicity of obtained
ophiostomatoid fungi to broaden the knowledge and
evaluate the potential threat of aforementioned fungal
isolates, which could be harmful to Scots pine.

**Material and methods.** Field study and sample
*collection.* In May 2017 and November 2018 to
examine the presence of ophiostomatoid fungi, samples
of blue-stained wood were taken from Scots pine trees
damaged by bark beetles during sanitary felling in Sumy
region: State Enterprise «Okhtyrske Forest Economy»
(Okhtyrske FE), Okhtyrske forestry, compartment 52,
subcompartment 3 and State Enterprise «Hostkinske
Forest Economy» (Hostkinske FE), Myronivske
forestry, compartment 5, subcompartment 27.

Sampling has been done from 28 randomly selected
pine trees (14 trees in each site) which were 58-102
years old showing external signs of crown dieback and
bark beetle attacks. Health condition (HC) of sampled
trees was evaluated visually before cutting on a range
of visual characteristics (crown color and density, dead
branches in the crown, etc.) according to «Sanitary rules
in the forests of Ukraine» (Sanitarni pravila v lesah
Ukrainy, 2016). Each tree was ranked according to one
out of six categories (1st – healthy; 2nd – weakened;
3rd – severely weakened; 4th – drying; 5th – recently
died; 6th – died over a year ago).

The two sites were approximately 256 km from
each other. Samples were taken manually and the tool
superficially sterilized with 96% ethanol after each
sampling. Samples of the bark beetle galleries, phloem,
and sapwood (approximately 5 × 5 cm) were stored in
separate clean paper bags in the fridge in the laboratory.

Samples were inspected in the laboratory using a
microscope for the presence of the fungal fruit-
* bodies in the bark beetle galleries. Wood samples
were sterilized by 96% ethanol for 15 seconds and
small wood fragments were used to place on selective
media for ophiostomatoid fungi (2% malt extract agar
with cycloheximide (200 mg/L, AldrichSigma, St.
Louis, Co. LLC.). Fungal cultures were purified by
transferring growing hyphae or spore masses from
individual colonies to new 2% malt extract agar
without cycloheximide. Cultures were incubated at 22 °C
in the dark. All cultures were grouped morphologically
and identified using identification keys. Due to
contradiction in the taxonomy of ophiostomatoid fungi,
molecular methods (Davydenko et al., 2017) were also
used to identify species.

**DNA extraction, PCR, sequencing, and molecular
identification.** DNA was extracted from the cultures
of the isolates representing morphological groups.

Approximate DNA concentrations were determined
at 260 nm using the Nano-drop 2000 spectrophotometer
(Nano-drop Technologies, Wilmington, DE, USA).

Internal transcribed spacer (ITS) regions 1 and 2,
including the ribosomal 5.8S gene, were amplified
using the ITS primer pairs (Davydenko et al., 2017).

The reaction mixture contained, in a total volume of
15 μl, 200 μM deoxyribonucleotide triphosphates,
0,2 μM of each primer, 0,03 U/μl Thermo Green Taq
polymerase with reaction buffer Green, and 2,75 mM
final concentration of MgCl2. The thermal cycling was
carried out using an Applied Biosystems GeneAmp PCR
System 2700 thermal cycler (Foster City, CA, USA).

An initial denaturation step at 95°C for 5 min was followed
by 35 amplification cycles of denaturation at 95°C
for 30 s, annealing at 55°C for 30 s, and extension at
72°C for 30 s. The thermal cycling was ended by a final
extension step at 72°C for 7 min. The protein-coding
gene β-tubulin (partial gene) was also amplified using
the primers Bt2a and Bt2b (Davydenko et al., 2017).

**Inoculation of pine seedlings.** To investigate the
pathogenicity of the ophiostomatoid fungi to
P. sylvestris, inoculation experiments were carried out on 3-5-year-old seedlings of P. sylvestris, growing outdoor of laboratory «Kharkivlisozahyst». Stem diameters at the inoculation point were between 12.6 mm and 14.8 mm (13.7 mm on average). All species from ophiostomatoid group were used in the pathogenicity test. In May 2019, 96 seedlings of P. sylvestris were inoculated with a spore suspension of selected fungal species (12 seedlings per fungus). All pine seedlings were inoculated by making a wound on the stem with the aid of a scalpel and pouring 10 μL of spore solution suspension (5×10³ spores mL⁻¹) onto the wound. The wound was then covered with a strip of Parafilm to prevent desiccation. In addition, 12 control seedlings were inoculated in the same way, with sterilized water, and the wounds were also covered with a strip of Parafilm. The morphological condition of all pine seedlings was first evaluated on a weekly, and then a monthly basis as visually healthy, weakened, or dead plants subject to typical symptoms of damping-off, crown decline, resin flow, and necrosis. Disease susceptibility was finally estimated at 120 and 180 days post-inoculation. After this, all plants were harvested and the bark of each seedling was peeled off the stem and the sizes of any lesions (length, depth, and width) that had formed in the phloem around the inoculation points were measured. The stems were then cut across the inoculation points. The tangential width and radial length (depth) of discoloured zones of sapwood were measured on the surface of the cross-section. Small pieces of wood were cut from the discoloured sapwood of the seedlings and transferred onto 2% malt extract agar with 0.8% cyclohexamide and 0.2% streptomycin sulphate (CSMA) to confirm the presence of the inoculated fungus.

Statistical analyses. Raw sequence data were analyzed using the SeqMan Pro version 10.0 software from DNASTAR package (DNASTAR, Madison, WI, USA). Databases at GenBank and at the Department of Forest Mycology and Plant Pathology, Swedish University of Agricultural Sciences, were used to determine the identity of ITS rRNA sequences. The criteria used for identification were: sequence coverage > 80%; similarity to taxon level 98-100%, similarity to genus level 92-97%.

A NCBI BLAST (National Centre for Biotechnology Information, www.ncbi.nlm.nih.gov) search was run with the edited sequences for preliminary species identification. Obtained data were analyzed using Kolmogorov–Smirnov test to check the normal distribution and homogeneity of variance. Data on lesion size of Scots pine seedlings inoculated with fungal species were analyzed using ANOVA general linear model (GLM) in Statistica software (STATISTICA® 7.0 (StatSoft, Inc., Tulsa, USA). Necroses were measured in only one direction from the point of inoculation. For ANOVA significant effects (p < 0.05), a post hoc HSD Tukey test was used to compare the means with a significance level of p < 0.05.

Results and discussion. Healthy, weakened, and dying Scots pine trees in both sites were inspected and visually evaluated to rank. The signs of bark beetle attack and typical symptoms of the presence of ophiostomatoid fungi (blue to greyish-black stains on wood) were observed only in trees of the 3rd–5th categories of health condition (severely weakened, drying, and recently died respectively) (Fig. 1).

Moreover, almost all trees in the 3rd–5th categories were damaged by bark beetles and ophiostomatoid fungi for both sites (Fig. 2).
In total, 256 isolates were obtained from wood samples which further were grouped in 8 clusters (Tab. 1). The isolates were used for DNA extractions, species identification, and further inoculation.

The most commonly for all trees at both sites was *Ophiostoma piceae* (82.1%), the slightly less abundant were *Ophiostoma ips* and *O. minus* (67.9%), as well as *Ophiostoma bicolor* (64.3%), (data not shown). The frequency of other detected species was lower than 30%. The five from eight blue-stain fungi (*Ophiostoma bicolor, O. ips, O. canum, O. piceae, O. minus*) were isolated from trees 3rd and 4th health categories at both sites. *Graphium* sp was found only on severely weakened trees at Shostkinske FE, while *Grosmannia* sp.1 was also isolated only from trees of 4th category of FHC but for both sites (see Tab. 1).

Table 1

| Ophiostomatoid species          | Frequency of isolates from galleries |
|---------------------------------|--------------------------------------|
|                                 | Okhtyrske FE | Shostkinske FE |
|-------|---------------|---------------|
|       | HCl=3 | HCl=4 | HCl=3 | HCl=4 |
| *Grosmannia* sp.1               | –       | 21.43 | –       | 35.71 |
| *Graphium* sp.                   | –       | –     | –       | 50.00 |
| *Ophiostoma bicolor* R.W. Davidson & D. E. Wells | 14.29 | 35.71 | 21.43 | 57.14 |
| *Ophiostoma canum* (Münch) Syd. & P. Syd | 7.14 | 14.29 | 14.29 | 28.57 |
| *Ophiostoma ips* (Rumbold) Nannfeldt | 28.57 | 28.57 | 28.57 | 50.00 |
| *Ophiostoma minus* (Hedgc.) Syd. & P. Syd | 28.57 | 14.29 | 35.71 | 57.14 |
| *Ophiostoma piceae* (Münch) Sydow & P. Sydow | 21.43 | 50.00 | 28.57 | 64.29 |
| *Ophiostoma* sp.1                | 14.29 | 28.57 | –       | –       |

The principal components analysis (PCA) based on Euclidean distances revealed that the first principal component explained 97.42% of the variability in the data of the fungal communities (data not shown). Four different clusters were formed for all sites in the respect to the fungal community compositions, indicating the significant difference between sites caused blue-stain fungal community shifts associated with bark beetles’ galleries and the similarity between the trees of the 3rd and 4th health condition categories (see Fig. 2).

The similarity matrices between tree categories and the difference between site groups were employed to yield different results. In this work, a similarity matrix based upon Euclidean distance, commonly used in cluster analysis, is developed as a viable alternative. This analysis resulted in Euclidean similarity-based PCA identifies parameters that are close to each other, thereby providing a wide range of similarity between fungal communities for trees 3rd and 4th HC measures available to investigators, to be chosen based on what characteristic they wish to identify. Moreover, the correspondence analysis revealed associations between fungal taxa and different types of samples (Fig. 3).
Ophiostoma ips and O. piceae were commonly associated with the pine trees from Okhtyrske FE while Ophiostoma bicolor and O. minus were commonly linked to the pine trees 3rd and 4th FHC from Shostkynse FE. Another group of fungi including Grosmannia and Ophiostoma canum were associated with all groups while Graphium sp. and Ophiostoma sp.1 were distributed more or less randomly and did not show a clear association with any type of sample (see Fig. 3).

Pathogenicity test. As mentioned in the introduction, some ophiostomatoid fungi are pathogenic to their host plants. Generally, ophiostomatoid fungi are introduced to their host plants through wounds produced by bark beetles feeding and gallery construction. The introduction and spread of ophiostomatoid fungi under the bark result in the necrotic lesions in the phloem around bark beetle galleries, sapwood colonization, and further, sometimes, tree decline or death. Therefore, artificial inoculation and further observation of emerged symptoms is the main criterion to determine the pathogen of particular plant disease. Therefore, the virulence of the ophiostomatoid fungi was evaluated by measuring the lesions and monitoring the mortality of Scots pine seedlings after inoculation for 6 months.

The inoculated seedlings were harvested after inoculation, and the size of the lesions formed in the inner bark around the inoculation points together with the sizes of the stained and dried zones of sapwood were measured. The necrotic lesions formed around the inoculation points on the inner bark spread mainly longitudinally and only slightly in the tangential direction. The lengths of lesions differ according to the fungus inoculated. Additionally, a wedge-shaped dry zone of sapwood with or without blue staining spreads from the inoculation point toward the border between the sapwood and heartwood. The fungi produced larger lesions and were more virulent. The ability of the fungi to colonize sapwood and disturb water transportation is important for evaluating their ability to kill host trees (Solheim, 1992; Yamaoka et al., 2000). Some ophiostomatoid fungi associated with bark beetles have the ability to kill their host trees; e.g., Endoconidiophora polonica associated with I. typographus (Solheim, 1992) and Grosmannia clavigera with Dendroctonus ponderosae (Yamaoka et al., 2000).

In our study, all isolated ophiostomatoid fungi were tested to check their ability to infect seedlings of *P. sylvestris*. Most of the pathogenicity results were generally concurred with previously described inoculation experiments, thereby demonstrating similar lesion morphology and similar patterns of plant tissue colonization by different ophiostomatoid fungal species (Jankowiak & Bilanski, 2013; Solheim, 1992). However, a few isolates, such as Graphium sp., Grosmannia sp., Ophiostoma sp., were tested first in our study. Ophiostomatoid fungi associated with *Ips acuminatus* and *Hylurgus ligniperda* infesting Scots pine in Ukraine were reported by Davydenko (2014, 2017). Revised information for these fungi together with unpublished data are shown in Table 2. Therefore, one species of Grosmannia and one species of Graphium were isolated from bark beetle galleries on infesting Scots pine.

According to our results, *O. minus* was much more pathogenic than other fungi based on all criteria of fungal pathogenicity. It was the only fungus that caused the highest rate of sapwood desiccation and mortality of 16.7% of seedlings in 6 months (see Tab. 2). *O. minus* caused dieback of *P. sylvestris* seedlings, which also demonstrated in previous studies (Solheim, 1992; Jankowiak, 2006). Most of the studies focused on pathogenicity ophiostomatoid fungi indicated that the ability to invade sapwood and phloem can be considered as critical for pathogenic colonization (Solheim, 1992; Linnakoski et al. 2012).

All eight fungal species inoculated caused resin exudation and staining on the bark around inoculations in Scots pine seedlings. The isolate of *O. minus* showed the highest growth in all directions, with statistical significance (*P* < 0.00147), although in the radial direction the growth was similar to *Graphium* sp. and in the tangential direction was similar to *O. bicolor* (see Tab. 2). This species produced a black discoloration of the wood and caused the greatest stained area (*P* < 0.001), as measured on cross-sections.

### Table 2

| Species                          | Phloem necrosis, mm** | Resin flow, % | Declining plants, % | Dead plants, % |
|----------------------------------|-----------------------|---------------|---------------------|---------------|
|                                  | Length | Width | Depth of blue-stain | 1 | 2 | 3 | 4 | 5 | 6 | 7 |                  |
| **Grosmannia sp.1**              |        |       |                     | 1 | 2 | 3 | 4 | 5 | 6 | 7 |                  |
|                                  | 13.24±0.22e           | 2.42±0.4b     | 1.29±0.4b            | 75.00 | 16.67 | 0.00 |
| **Graphium sp.**                 |        |       |                     | 1 | 2 | 3 | 4 | 5 | 6 | 7 |                  |
|                                  | 19.86±0.12d           | 3.24±0.6c     | 1.62±0.4b            | 33.33 | 41.67 | 0.00 |
| **Ophiostoma ips** (Rumbold)     |        |       |                     | 1 | 2 | 3 | 4 | 5 | 6 | 7 |                  |
| **Nannfeldt**                    | 4.52±0.15c            | 2.58±0.2b     | 0.21±0.2a            | 25.00 | 0.00  | 0.00 |
|                                  | 42.56±0.1d            | 6.62±1.1c     | 1.65±0.4b            | 91.67 | 41.67 | 16.67 |
The second fastest-growing fungal species in all directions was Graphium sp. This species caused a grey stain on the wood and decline 41.7% of pine seedlings as well as O. bicolour (see Tab. 2). Grosmannia sp.1 grew at a similar radial rate to Graphium sp. but grew more slowly both tangentially and especially longitudinally. Ophiostoma sp.1 caused the decline of 33% pine seedlings and grew significantly (P < 0.001) more slowly than Graphium sp. Ophiostoma canum and O. piceae caused significant less decline of pine seedlings and grew much more slowly than above-mentioned ophiostoma fungi (16.67 and 8.33% respectively, P < 0.001) while O. bicolour and O. ips did not caused seedling decline at all during 6 months (see Tab. 2), although caused staining on the bark around inoculations in all directions significantly more than control inoculations (P < 0.001). O. ips and O. piceae penetrated the smallest areas of the wood caused a light greyish discoloration compared to the other species (P < 0.05).

The inoculated fungi were successfully re-isolated from 98% of the inoculation points. No ophiostomatoid species were isolated from the control trees.

**Conclusions.** Based on the ability of tree declining, killing and wood staining, *O. minus* is the most important species on Scots pine trees, followed by *Graphium* sp. and *Ophiostoma* sp.1. The ophiostomatoid fungi isolated in this study are commonly found to be associated with bark beetles and detected on conifer timber (Linnakoski et al., 2012). The occurrence of ophiostomatoid species in the sapwood of Scots pine is consistent with the concept the ophiostomatoid fungi are primary colonizers of the fresh sapwood of trees and timber and other wood in the succession of microorganisms during wood decay (Solheim, 1992).

### Список літератури

Санітарні правила в лісах України (2016); у редакції постанови Кабінету Міністрів України від 26 жовтня 2016 р. № 756 (Sanitary Forests Regulations in Ukraine (2016)). In the redaction of Decree of Cabinet Minister of Ukraine from 26 October 2016, № 756 (in Ukrainian)). Retrieved from https://zakon.rada.gov.ua/laws/show/555-95-n

Buano, A., Diez, J. J., & Fernandez, M. M. (2010). Ophiostomatoid Fungi Transported by *Ips sexdentatus* (Coleoptera, Scolytidae) in Pinus pinaster in NW Spain. *Silva Fennica*, 44 (3), 387-397. https://doi.org/10.14214/sf.137

Cottrell, S., Mattor, K. M., Morris, J. L., Fettle, C. J., McGrady, P., Maguire, D., …Roberts, R. (2020). Adaptive capacity in social-ecological systems: a framework for addressing bark beetle disturbances in natural resource management. *Sustainability Science*, 15 (2), 555-567. https://doi.org/10.1007/s11625-019-00736-2

Davydenko, K. (2019). A comparative characteristic of fungal communities associated with *Ips acuminatus* in different regions of Ukraine. *Наукові праці Лісівничої академії наук України*, 18, 118-128. https://doi.org/10.15421/411912

Davydenko, K., Vasaitis, R., & Menkis, A. (2017). Fungi associated with *Ips acuminatus* (Coleoptera: Curculionidae) in Ukraine with a special emphasis on the pathogenicity of ophiostomatoid species. *European Journal of Entomology*, 114, 77-85. https://doi.org/10.14411/eje.2017.011

Davydenko, K., Vasaitis, R., Meshkova, V., & Menkis, A. (2014). Fungi associated with the red-haired bark beetle, *Hylurgus ligniperda* (Coleoptera: Curculionidae) in the forest-steppe zone in eastern Ukraine. *European Journal of Entomology*, 111 (4), 561-565. Retrieved from https://search.proquest.com/docview/1622264211?accountid=28676

De Beer, Z. W., Seifert, K., & Wingfield, M. (2013). A nomenclator for ophiostomatoid genera and species in the Ophiostomatales and Microascales. *The ophiostomatoid fungi: expanding frontiers*, 12, 245-322. Retrieved from https://www.fungaltaxonomy.org
Офіостомові гриби, які пов’язані з короїдами та заселяютьPinus sylvestris у Сумській області (Україна)

К.В. Давиденко1, Д.О. Батуркін2

Останнім часом у східному регіоні України, зокрема в Сумській обл., відбувається інтенсивне виявляння непов’язаних з оселінними лісами гімнофітів короїдів іх насадження. Короїд губить насадження всередині, які пошкоджують два види короїдів – Ips acuminatus і Ips sexdentatus. Офіостомові гриби, які переносять короїді, можуть значно посилювати загальну шкоду лісам від цих комах, що необхідно брати до уваги під час оцінювання загальних збитків від короїдів.

Метою дослідження було виявити видовий склад офіостомових грибів, пов’язаних із сильно пошкодженими деревами сосни звичайної у Сумській обл., оцінити патогенності офіостомових грибів і встановити потенційну загрозу для Pinus sylvestris L.

Офіостомові гриби виділені з личинкових ходів короїдів та ідентифіковані до рівня виду або роду на основі морфологічних особливостей та аналізу ITS послідовностей ДНК.

Галузь виділено вісім видів офіостомових грибів (Graphium sp., Grosmannia sp.1, Ophiostoma bicolor, O. ips, O. canum, O. piceae, O. minus, Ophiostoma sp.1). Саджанці Pinus sylvestris іонкувані Wingfield, M.J., Barnes, I., de Beer, Z.W., Roux, J., Wingfield, B.D., & Taurum, S.J. (2017). Novel associations between ophiostomatoxiphi fungi, insects and tree hosts: current status-future prospects. Biological Invasions, 19 (11), 3215-3228. https://doi.org/10.1007/s10530-017-1468-3

Yamaoka, Y., Takahashi, I., & Iguchi K. (2000). Virulence of ophiostomatoxiphi fungi associated with the spruce bark beetle Ips typographus f. japonicus in Yezo spruce. Journal of Forest Research, 5, 87-94. https://doi.org/10.1007/BF02762525

1 Давиденко Катерина Валеріївна – член-кореспондент Лісівничої академії наук України, кандидат сільськогосподарських наук, старший науковий співробітник. Український науково-дослідний інститут лісового господарства та агролісомеліорації ім. Г.М. Висоцького, вул. Пушкінська, 86, Харків, 61024, Україна; Запрошений науковий співробітник Департаменту Лісової мікології і фітопатології Шведського Аграрного Університету, P.O. Box 7026, SE-75007, Уппсала, Швеція. Тел.: +38-098 66 755 26. E-mail: kateryna.davydenko74@gmail.com ORCID http://orcid.org/0000-0001-6077-8533

2 Батуркін Денис Олександрович – директор Державного спеціалізованого лісозахисного підприємства «Харківські ліси», Харківська обл., смт. Покотилівка, вул. Незалежності, 127, 62458, Україна; Запрошений науковий співробітник Лісівничої академії наук України, кандидат сільськогосподарських наук, старший науковий співробітник. Український науково-дослідний інститут лісового господарства та агролісомеліорації ім. Г.М. Висоцького, вул. Пушкінська, 86, Харків, 61024, Україна. Тел.: +38-067 764 81 07. E-mail: baturkin.denis@ukr.net ORCID https://orcid.org/0000-0002-6061-9863
Целью исследования было выявить видовой состав офиостомовых грибов, связанных с сильно ослабленными и усыхающими деревьями сосны в Сумской обл., оценить патогенность офиостомовых грибов и потенциальную угрозу для Pinus sylvestris.

Офиостомовые грибы выделены из личиночных ходов короедов и определены до уровня вида или рода с помощью микроскопа по морфологическим особенностям, а также с помощью молекулярного метода, используя анализ ITS последовательностей ДНК.

Всего выделено восемь видов офиостомовых грибов (Graphium sp., Grossmannia sp. 1, Ophiostoma bicolor, O. ips, O. canum, O. piceae, O. minus, Ophiostoma sp. 1). Саженцы Pinus sylvestris инокулированы восемью видами выделенных грибов и стерильной средой в качестве контроля для оценки их патогенности. Инокулированные растения оставляли в течение шести месяцев, оценивая общее состояние растений, размер некрозов, потеки смолы. Инокуляция саженцев грибом O. minus привела к наиболее интенсивным поражениям и даже гибели сосновых саженцев. Все восемь видов грибов вызвали экссудацию смолы и обесцвечивание коры вокруг места инокуляции. Пять видов грибов вызвали ослабление саженцев в той или иной степени. Площадь обесцвечивания флоэмы также была достоверно большей у растений, инокулированных грибом O. minus. Все офиостомовые грибы вызывали некрозы древесины большего размера по сравнению с контролем. По результатам исследования, в связи со способностью вызывать усыхание и гибель растений, офиостомовые грибы, связанные с сосной, следует считать одним из опасных патогеном сосновых саженцев. Менее вирулентными видами оказались Graphium sp. и Ophiostoma sp. 1. Факт заселения усыхающих деревьев сосны офиостомовыми грибами согласуется с концепцией, что они являются первичными колонизаторами свежей древесины в процессе ее разрушения.

**Ключевые слова:** офиостомовые грибы; сосна обыкновенная; короеды; патогенез; санитарный статус.