Preliminary results on narrow-leaved ash (Fraxinus angustifolia Vahl) and green ash (Fraxinus pennsylvanica Marshall) seed entomofauna in Croatia

Jelena Kranjec Orlović, Ida Bulovec*, Milivoj Franjević, Damjan Franjević, Josip Skejo, Marin Biliškov, Danko Diminić, Boris Hrašovec

Summary

Native narrow-leaved ash (Fraxinus angustifolia Vahl) and introduced green ash (Fraxinus pennsylvanica Marshall) are two most common ash (Fraxinus spp.) species in Croatian lowland forests, taking a significant place in these ecosystems. In the recent past it has been observed that narrow-leaved ash seed yields have reduced, emphasizing the importance of good health status of those collected, as they are used for seedling production and consecutive regeneration of forest stands or afforestation. Given that insects could be an important factor affecting ash seed health status, the aim of this research was to gain insight into the species being present and their frequency in the ash seeds. In total 2.500 narrow-leaved ash and 1.000 green ash seeds from seven different locations were screened for insect presence. Analysed seeds were categorized as undamaged, insect damaged or empty, and observed insect specimens were identified by morphological and molecular genetic analysis. Green ash seeds revealed higher insect infestation in comparison with native ash species, with weevil Lygniodes bischoffii being the most numerous pest found. Narrow-leaved ash seeds were mostly occupied by the native weevil species, L. enucleator. Besides these pests, larval, pupal, and adult stages of parasitoid wasps (Hymenoptera) and larval stage of the seed eating moth Pseudargyrotoza conwagana (Lepidoptera) were found in both ash species. However, given that pests detected in this study prefer urban areas or were present in a very small number of seeds, where they were often followed by parasitoids, it can be concluded that insects currently do not have a significant negative impact on the reproduction of narrow-leaved ash and green ash in forest stands.

Key words: ash, seed damage, weevils, Lygniodes, parasitoid wasps, Pseudargyrotoza conwagana
INTRODUCTION

Narrow-leaved ash (Fraxinus angustifolia Vahl), distributed in central-southern Europe and northwest Africa, is an ecologically and economically important species in riparian and floodplain forests because of its fast growth and specific adjustments to the habitat, especially in central Europe, the Pannonian Basin and the Balkans (Temunović et al. 2012, Caudullo and Durrant 2016). It is currently suffering severe dieback in its distribution area (Gross et al. 2014), being one of the most endangered forest tree species in Croatia, mostly due to a pathogenic fungus Hymenoscyphus fraxineus (T. Kowalski) Baral, Queloz & Hosoya (Diminić 2015, Milotić et al. 2016). Yield and health status of narrow-leaved ash seeds are thus of great significance for seedling production and consecutive afforestation or regeneration of damaged forest stands. Previous studies showed that freshly collected seeds have a very high viability even after two years of storage (87%) and are of relatively good health status. Collected seeds have a very high viability even after two years of storage (87%) and are of relatively good health status. Generally, most studies focusing on entomofauna in the ash seeds in Europe and their impact on seed health status were conducted on common ash (Fraxinus excelsior L.) (e.g. Gardner 1977, Tapper 1992, Hayatgheibi 2013, Gosik et al. 2017), and there is little or no information about insect presence in seeds of other ash species. Therefore, this research had two main objectives: (1) to explore entomofauna of narrow-leaved ash and green ash seeds in order to gain insight into the insect species present in the most common native and introduced ash species in Croatia for future comparisons; (2) to show the frequency in which insects appear in seeds, as this could be an important factor affecting ash seed health status.

MATERIALS AND METHODS

Narrow-leaved ash seeds were collected in the period from August to November 2017 from visually healthy trees in natural or specially managed forest stands registered as seed sources or seed stands (five different locations). Additional seed sampling was conducted from August to September 2018 in natural forest stands registered as the seed source HR-FAN-SI-121/305. Green ash seeds were collected in November 2018 from two different locations, one in an urban area and another in a natural forest stand (Figure 1).

One hundred seeds from each location were separated from samaras and examined for presence of insects under the stereo microscope (SMZ 168-TLED, Motic, Hong Kong, China). The seeds were categorized as undamaged (without any sign of insect presence), damaged (with presence of different life stages of insects or their exit holes), or empty (partially developed but with no endosperm). Collected insect specimens were analysed and photographed under the stereo microscope and grouped according to their morpho-
logical characteristics into presumed insect taxa, which were preserved in 75% (v/v) ethanol solution, each in a separate microcentrifuge tube. As the number of individuals in sampled seeds was generally low, additional 2,000 of narrow-leaved ash seeds from Županja and 800 green ash seeds from Zagreb and Sunja were examined in order to collect as much biological material as possible for species identification by molecular genetic analysis.

Total genomic DNA was extracted from ethanol-preserved specimens with DNeasy Blood & Tissue kit (Qiagen, Hilden, Germany) according to the manufacturer’s instructions. The Folmer region of mitochondrial CO1 gene (cytochrome c oxidase 1) was amplified from obtained DNA using primers LCO1490 and HCO2198 (Folmer et al. 1994), in 50 µL PCR reactions containing 25 µL TopTaq Master Mix kit (Qiagen, Hilden, Germany), 1 µL of DNA, 1 µL of 10 µmol/L of each primer, and 22 µL of distilled RNA-free water. Cycling conditions were as follows: an initial denaturation at 95 °C for 5 min, 35 cycles of denaturation at 94 °C for 1 min, annealing at 54 °C for 1 min, extension at 72 °C for 1 min and a final extension step at 72 °C for 10 min. The resulting PCR products were purified with MinElute PCR Purification kit (Qiagen, Hilden, Germany) according to the manufacturer’s protocol and sequenced at the DNA sequencing facility (Macrogen Europe, Amsterdam, The Netherlands) using PCR amplification primers. The obtained sequences were identified by comparison with reference sequences in NCBI GenBank (Benson et al. 2017) and BoLD (Ratnasingham and Hebert 2007) databases. Species were identified using both morphological and molecular data.

RESULTS

Inspection of 100 seeds per location revealed that 10% or less of narrow-leaved ash seeds were damaged by insects from all locations except Vukovar, whereas more than third of green ash seeds from both locations were occupied by insects (Figure 2). In total, 10.6% of narrow-leaved ash seeds and 42% of green ash seeds displayed insect damage or presence.

Damaged seeds of both ash species contained weevil larvae (Coleoptera: Curculionidae), wasp larvae and adults (Hymenoptera), and moth larvae (Lepidoptera), while wasp pupae and fly larvae (Diptera) where found only in green

![Figure 1. Map of Fraxinus angustifolia and Fraxinus pennsylvanica seed collection locations](image1.png)

![Figure 2. Proportion of undamaged, insect damaged and empty Fraxinus angustifolia and Fraxinus pennsylvanica seeds from seven different locations](image2.png)
ash seeds (Table 1). Weevil and moth larvae found in narrow-leaved ash seeds were identified based on the morphological and molecular genetic analysis, as *L. enucleator* and *Pseudargyrotoza conwagana* (Fabricius, 1775, Lepidoptera: Tortricidae), with 98.21% and 99.06% sequence similarity to reference sequences in NCBI GenBank, respectively. Wasp larvae and adults found in this ash species were not successfully identified (Figure 3). Morphological and molecular genetic analysis of wasps in damaged seeds of green ash revealed that found pupae belonged to species *Eupelmus urozonus* (Dalman, 1820, Hymenoptera: Eupelmidae), and larvae to family Pteromalidae (Hymenoptera), with 99.35% and 93.30% sequence similarity to reference sequences in NCBI GenBank, respectively. Wasp adults remained unidentified. Weevil larvae were identified as *L. bischoffi*, combining their morphological characteristics according to Gosik et al. (2017) and molecular genetic analysis, because the obtained sequence was ambiguous and shared only 89.53% similarity with other *L. bischoffi* sequences deposited in NCBI GenBank. Moth larvae was identified based on its morphological characteristics as *P. conwagana*. Fly larvae were not successfully identified (Figure 3).

**DISCUSSION AND CONCLUSIONS**

**RASPRAVA I ZAKLJUČCI**

Relatively small number of damaged and empty narrow-leaved ash seeds was found in this research, implying their good health status regarding insect infestation. Although seeds of introduced green ash revealed higher insect occupation, especially those collected in the urban area, most of the specimens found (81%) were parasitoids, indicating that pest populations are naturally held under control. In total, four insect taxa were identified to species level in this research, of which three can be categorised as seed pests (*L. enucleator*, *P. conwagana*, and *E. urozonus*).
enucleator, L. bischoffi, P. conwagana). Other specimens were mostly parasitoid wasps in different stages, which, with the exception of E. urozonus pupae, remained unidentified.

L. enucleator, found only in native narrow-leaved ash seeds in this research, is the most common and widespread species of the Palearctic Lignyodes (Clark and Lodos 1981), present in the most part of the Europe on ash seeds and seeds of Ligustrum vulgare L. and Syringa vulgaris L. (Dieckmann 1970, Mazur 2002, Arndt and Hielser 2007, Nakladal 2011, Bacal et al. 2013, Caldara 2013, Delbol 2013). Proportion of seeds infested by L. enucleator in the present study was only 4.64% altogether, which could be a consequence of its preference towards trees that are well exposed to the sun, mostly in rural and urban biotopes (Gosik et al. 2017), and avoidance of trees growing in riparian forests, where its larvae and pupae overwintering in the soil could be inundated and killed during spring floods (Kania et al. 2001; Gosik et al. 2001). The fact that analysed seeds were collected in regularly flooded narrow-leaved ash forest stands is the most probable cause for low rate of L. enucleator presence in seeds. Therefore, it is plausible to conclude that this weevil does not represent danger to the seed health status and reproduction rate of narrow-leaved ash in its natural habitat in Croatia.

In the present study, L. bischoffi was found only in the seeds of introduced green ash. In Europe this weevil is so far confirmed in Austria, Hungary, Moldova, Poland, Russia, Slovakia, Switzerland and Ukraine, most frequently on the North American species such as F. americana L., F. nigra Marsh and F. pennsylvanica, but also on F. ornus L., F. lanceolata Borkh., F. oxycarpa Will., and F. excelsior (Dieckmann 1974, Clark 1980, Caldara 2013, Bacal et al. 2013, Arzanov 2013, Gosik et al. 2017). This is the first official report of L. bischoffi in Croatia. Although the proportion of seeds infested by living larvae was only 2.1% in this study, in every seed where wasp larva was found there was a dead larva of the weevil as well; and since the ash seeds were collected at the beginning of November, some larvae had already left the seeds, leaving an exit hole. Combining those findings, the actual proportion of infested seeds was around 16%, but was significantly decreased by the present parasitoids. The fact that L. bischoffi prefers urban areas and trees surrounded by grassy areas well exposed to the sun...
Lithuania (Ostrauskas 2004) and Poland (Kubasik 2011). The other pest found on both ash species in this research was *P. conwagana*, a phytophagous moth from Lepidoptera family widespread through the Eurasia region on the plants belonging to genera *Ligustrum*, *Fraxinus*, *Syringa* and *Berberis* (Tapper 1992). In Europe large numbers of its caterpillars were reported in seeds of common ash in England, causing 15-36% seed loss (Gardner 1977), and also in Sweden (Tapper 1992, Hayatgheibi 2013), Slovakia (Kollar 2007), Lithuania (Ostrauskas 2004) and Poland (Kubasik 2011). There are no reports of this pest in narrow-leaved ash and green ash seeds generally, nor was it found in Croatia before. Currently, *P. conwagana* is not a threat to the examined ash species in Croatia, since it was found in only two narrow-leaved ash seeds from Županja and in one green ash seed from Sunja (both in woodland areas). Regarding narrow-leaved ash, the low infestation rate could be the consequence of a decrease in seed production (Vincenc 2015, 2016, 2017, 2018, 2019), as Janzen (1971) reports that natural fluctuations of ash species in seed production lead to lower rates of moth infestation. However, this explanation is not applicable to green ash, and it might be that in Croatia ash species are not the primary source of food for this seed-feeding tortricid, but rather some other plant species. In the seeds of both species of ash, parasitoid wasps were detected in the larval, pupal, and adult stages. The only determined species was *E. urozonus* collected from the green ash seeds from both sampling locations. This wasp is considered to be exceptionally polyphagous, as it occurs on hosts belonging to Lepidoptera (Gelechiidae, Oecophoridae, Tortricidae), Coleoptera (Bruchidae, Curculionidae, Scolytidae), Diptera (Agromyzidae, Cecidiomyiidae, Tephritidae), Neuroptera (Chrysopidae) and Hymenoptera (Braconidae, Tenthredinidae, Ichneumonidae) (Gibson and Fusu 2016). Although *E. urozonus* is reported to be established throughout Europe (Gibson and Fusu 2016), this is the first official report of its presence in Croatia. Comparing found specimens of adult wasps with the descriptions in the literature (Al khatib et al. 2014), there is a possibility that some of the acquired adults could also be *E. urozonus*, but this requires further investigation. Larval stages of parasitoid wasps found in green ash where determined to belong to Pteromalidae family, which is the largest family within superfamily Chalcidoidea with over 3,500 mostly parasitic species, often difficult to determine (Burks 2009). Wasp larvae in narrow-leaved ash and adults found in both species of ash could not be identified by molecular genetic analysis due to their small numbers and size, but were identified by their morphological characteristics as parasitoid wasps. For further determination they require a more detailed examination.

Comparing the health status of the seeds of the native narrow-leaved ash and the introduced green ash, the native seeds were generally in a better state. As for green ash seeds, those collected from a forest stand (Sunja) were less damaged by insects than the ones collected from the urban park in Zagreb, which is in accordance with the claim that *L. bischoffi* most often occurs in the urban areas (Wanat and Gosik 2003, Gosik 2006). Considering that two most numerous pests found in this study, *L. enucleator* and *L. bischoffi* prefer urban areas, that relatively large number of parasitoids was found in the seeds and on their larvae, and that other pests were present in a very small number of seeds, it can be concluded that insects currently do not have a significant negative impact on the reproduction of narrow-leaved ash and green ash in forest stands.

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SAŽETAK

Autohtoni poljski jasan (*Fraxinus angustifolia* Vahl) i alohtoni pensilvanski jasan (*F. pennsylvanica* Marshall) su dvije najčešće vrste roda *Fraxinus* prisutne u nizinskih šumama Hrvatske, zbog čega uzimaju važno mjesto u ovim šumskim ekosustavima. Posljednjih godina je zapaženo smanjenje uroda sjemena poljskog jasena, naglašavajući važnost dobrog zdravstvenog stanja sakupljenih količina, jer se sjeme koristi za proizvodnju sadnica koje se dalje upotrebljavaju za pomlađivanje šumske sastojina ili pošumljavanje. Kako kukci mogu imati velik utjecaj na zdravstveno stanje sjemena, cilj ovog istraživanja je bio dobiti uvid u prisutne vrste i njihovu učestalost u sjemenu jasena. Prisustvo kukaca je analizirano u ukupno 2.500 sjemenki poljskoga i 1.000 sjemenki pensilvanskog jasena. Promatrano sjeme je kategorizirano kao neoštećeno, oštećeno i prazno (šturo), a nađene jedinke kukaca su identificirane na temelju morfoloških karakteristika i korištenjem molekularno genetičke analize. Sjeme pensilvanskog jasena pokazalo je veću oštećenost kukcima u usporedbi sa sjemenu poljskoga jasena, pri čemu je pipa *Lygniodes bischoffii* bila najčešći utvrđeni štetnik. Oštećeno sjeme poljskog jasena je najčešće bilo nastanjeno autohtonom vrstom pipe, *L. enucleator*. Osim navedenih štetnika, u sjemenu obje vrste jasena su nađene ličinke, kukuljice i imaga parazitoidnih osa (Hymenoptera) te ličinke štetnika *Pseudargyrotoza conwagana* (Lepidoptera). S obzirom da štetnici utvrđeni u istraživanju preferiraju urbana područja ili su nađeni na vrlo malom broju sjemena, često zajedno sa parazitoīdimama, može se zaključiti kako kukci trenutno nemaju značajan negativan utjecaj na zdravstveno stanje sjemena poljskog i pensilvanskog jasena u šumskim sastojinama.

KLJUČNE RIJEČI: jasan, oštećenost sjemena, pipe, *Lygniodes*, parazitoidne ose, *Pseudargyrotoza conwagana*