Mites of the order Mesostigmata on plots reforested with Scots pine after various kinds of soil preparation and wood residue utilization after clear-cutting

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Abstract: The clear-cutting system is still popular, especially on poor sites in Central European forests. It causes serious disturbances in natural processes during forest regeneration. The main objective of this research on mesostigmatid mites was to evaluate and compare the influence of various kinds of soil preparation and treatment of the wood debris left after clear-cutting (on plots reforested with Scots pine *Pinus sylvestris*). As a result of the conducted investigations, 24 mite species of the order Mesostigmata were found. The highest number of species represented the families Parasitidae (7 species), Laelapidae (4 species) and Zerconidae, Digamasellidae, Ascidae (3 species each). The study was conducted on an experimental plot in the Międzychód Forest District (West Poland). The species composition of mite communities on all subplots is characteristic for Scots pine forests. The highest numbers of individuals and species were collected on the plots where the wood residue was burnt or chipped and left on the soil surface. The lowest numbers of individuals and species were collected on the plots where the chipped wood residue was mixed with soil or removed from the plots. The method of wood residue utilization significantly influenced the community structure of the Mesostigmata, while various kinds of soil preparation before reforestation did not have any significant influence.

Key words: Scots pine, Pinus sylvestris, reforestation, mites, Acari, Mesostigmata, species composition

INTRODUCTION

Forest management in Poland and in Europe is based on the principle of sustainability, formulated in 1804 by L.G. Hartig. According to this principle, present generations of people should maintain the forests in the way that the benefits gained from the forests by future generations will not be lower than those currently achieved (GORNOWICZ 2005). Sustainability may be achieved only on the basis of ecology. The proper development of forests requires the cycling of nutrients and a continuous flow of energy between the trophic levels of a forest ecosystem. Therefore, one has to establish what benefits may be gained from a forest so that the habitat will not suffer (e.g. due to soil acidification or loss of nutrients). A good way to maintain forest soil fertility is to leave wood debris (residue) at clear-cut sites, as the residue contains many minerals, which return to the soil, facilitate the decay of duff, and prevent the creation of acid humus (GORNOWICZ 2005). It is therefore worthwhile to find out which wood residue utilization methods have a positive influence on the soil, and the qualities of clear-cut sites. The soil fauna seems to be a good indicator of fertility and other natural qualities, as such organisms react to the changing environment relatively fast.

This research aimed at establishing which method of wood residue utilization and soil preparation provides best conditions for the soil fauna after a clear-cut and artificial reforestation, being a huge change in the local environment.

MATERIALS AND METHODS

The experimental plot of the Department of Forest Technology is situated in the Międzychód Forest Division (area 194d). It lies in the Noteć Forest (Puszcza Notecka), which is the largest compact forest complex in the Wielkopolska Lowland and the second largest in Poland (1200 km²). The forest grows on inland dunes (20-30 m high). Taking into account the Polish climate in general, the local climate is relatively warm and dry, with an average precipitation of 535–570 mm (KUSIAK & DYMEK-KU-SIAK 2002) and annual average temperature of 7.8–8.1°C (Woś 1994).

On the 3.3-ha plot, Scots pine *Pinus sylvestris* L. is grown on a potential site of fresh coniferous forest (labelled by Polish foresters as Bśw), on a deluvial soil formed on loose dune sand. In the spring of 2001, Scots pine (80%) and birch *Betula pendula* L. (20%) was planted in the plot, in the form of strips and clumps. In addition, grey alder *Alnus incana* (L.) Moench was planted between rows to improve the productivity of the soil. Grey alder is known to enrich the soil with nitrogen thanks to actinomycetes (symbiotic bacteria from the order Actinomycetales, now classified as the type Actinobacteria). This is especially advantageous in the poor soil of the plot.

The plot was divided into 3 belts (18 m wide, 600 m long), each with a different soil preparation method (A, B and C), in the north-south direction:

A = furrows with a ditch (the eastern belt, adjoining a young forest stand): furrows were ploughed with an active single-disc forest plough U-162 with a subsoiler, fixed to a tractor);

B = no preparation (the central belt);

C = classic furrows (the western belt, adjoining a 100-year-old forest): the soil was prepared by ploughing furrows with a double-mouldboard plough Lpz-75 with a ripper, used most often in forest sites.

The 3 differently prepared belts (basic belts) were divided by 5 transverse belts into 5 subplots in each basic belt. The transverse belts were subjected to various methods of wood residue utilization: 1 = raked and removed; 2 = burnt; 3 = left

undisturbed; 4 = chipped and left on the surface; 5 = chipped and mixed with the soil.

Soil samples from each subplot were taken 3 times in November 2005 with a sampler of 40 cm² (in total: 15 subplots and 45 soil samples). The samples were taken between trees, to the depth of 5 cm of mineral soil, together with litter and humus. The mites were extracted in Tullgren funnels. Permanent and fresh preparations were made.

The analysis of the results was conducted with the aid of the AnalizaTOR 2.0 software (Desmodus, Poland), using typical indices:

• species similarity between microhabitats (S), calculated as S = 2C/(A + B), where A = number of species in one microhabitat; B = number of species in another microhabitat; and C = number of species common to both microhabitats;

• dominance (*D*, in %), calculated as $\overline{D} = s/S$, where: s = number of individuals of a species; S = number of individuals of all species recorded in the habitat (ODUM 1982).

The following dominance classes were distinguished: eudominants (> 10%), dominants (5.1–10%), subdominants (2.1–5.0%), recedents (1.1–2.0%), subrecedents (<1.1%) (NIEDBALA et al. 1981). Also a multivariate analysis of variance (MANOVA) was conducted with the aid of SAS 9.1.3 software, to establish whether wood residue utilization and soil preparation methods have a significant influence on the number of mite individuals and species per sample.

The species names and taxonomy of Mesostigmata are consistent with BLASZAK & MADEJ (1997) and WIŚNIEWSKI (1997).

RESULTS AND DISCUSSION

A total of 1724 mites from the order Mesostigmata were collected. They represent 26 mite taxa of 10 families, of which 21 were identified to the species level and the others were identified to the genus level. The family Parasitidae was represented by 6 species, the Laelapidae by 4, while the Zerconidae, Digamasellidae and Ascidae by 3 species each. The species composition in all subplots was typical for pine forests growing on fresh coniferous forest sites (SKORUPSKI & BŁASZCZAK 1998). The list of Mesostigmata recorded in each of the differently prepared basic belts and their dominance indices are presented in Table 1. In all the differently prepared basic belts, eudominants include Paragamasus conus (Karg, 1971) and Gamasellodes bicolor (Berlese, 1918), whereas dominants are Zercon zelawaiensis (Sellnick, 1944) (only on belt A as a eudominant) and Asca aphidiodes (Linnaeus, 1758) (only on belt A as a recedent), Paragamasus vagabundus (Karg, 1968) (on belt C) and Amblyseius sp. (on belt A). The list of Mesostigmata mites recorded on transverse belts (different wood residue utilization methods), and their dominance indices are presented in Table 2. The dominant species in all transverse belts are Paragamasus conus and Gamasellodes bicolor, but their dominance in individual subplots varied widely, from 12.3 to 40.54 for Paragamasus conus and from 22.58 to 60.43 for Gamasellodes bicolor. Eudominants included also Paragamasus vagabundus (belt 1) and Zercon zelawaiensis

	Dominance index (%)			
Species	belt A	belt B	belt C	
Amblyseius sp.	8.11	2.53	3.35	
Anthoseius sp.	0	0	0.15	
Asca aphidioides (Linnaeus, 1758)	1.69	6.53	5.78	
Asca bicornis (Canestrini & Fanzago, 1887)	1.52	0	4.87	
Dendrolaelaps foveolatus (Leitner, 1949)	0.34	0	0	
Dendrolaelaps latior (Leitner, 1949)	2.36	3.16	0.91	
Dendrolaelasp sp.	1.01	0	0.15	
Gamasellodes bicolor (Berlese, 1918)	38.68	29.26	31.2	
Hypoaspis aculeifer (Canestrini, 1883)	1.01	2.32	0.76	
Hypoaspis karawaiewi Berlese, 1903	1.35	1.89	0	
Hypoaspis lasiomyrmecophilus Hirschmann, 1969	0	0	0.46	
Hypoaspis praesternalis Willmann, 1949	0.17	1.68	0.15	
Leptogamasus tectygynellus Athias-Henriot, 1967	0.84	4.63	1.83	
Paragamasus conus (Karg, 1971)	25.34	30.11	34.4	
Paragamasus sp.	0	0	0.3	
Paragamasus vagabundus (Karg, 1968)	1.35	3.58	5.78	
Parasitidae sp.	0	0.21	0.3	
Parazercon radiatus (Berlese, 1914)	0	0.21	0.61	
Pergamasus crassipes (Linnaeus, 1758)	0	0.63	1.37	
Pergamasus sp.	0	0	0.15	
Rhodacarus mandibularosimilis Shcherbak et Kadite, 1980	1.01	2.74	0.76	
Trachytes aegrota (C.L.Koch, 1841)	0.17	0.21	0	
Trichouropoda ovalis (C.L. Koch, 1839)	0.17	1.47	0.15	
Veigaia cervus (Kramer, 1876)	0	0.21	0.15	
Veigaia nemorensis (C.L. Koch, 1839)	0.84	1.89	0.3	
Zercon peltatus C.L. Koch, 1836	0	0.42	0	
Zercon zelawaiensis Sellnick, 1944	14.02	6.32	6.09	

Table 1. Dominance index (%) of mesostigmatid mites found in differently prepared basic belts (A = furrow with a ditch; B = no preparation; C = classic furrow)

		Dominance index (%)				
Species	belt 1	belt 2	belt 3	belt 4	belt 5	
Amblyseius sp.	5.35	3.29	0.79	7.94	6.01	
Anthoseius sp.	0	0	0	0	0.3	
Asca aphidioides	0	2.74	3.54	8.93	5.71	
Asca bicornis	0	6.03	1.57	0.99	0	
Dendrolaelaps foveolatus	0.53	0.18	0	0	0	
Dendrolaelaps latior	1.07	4.02	1.97	1.49	0	
Dendrolaelasp sp	0	0.18	2.36	0	0	
Gamasellodes bicolor	60.43	38.57	24.8	22.58	28.53	
Hypoaspis aculeifer	0.53	0.73	3.15	1.24	1.2	
Hypoaspis karawaiewi	0	0.37	2.36	0.5	2.1	
Hypoaspis lasiomermecophilus	1.6	0	0	0	0	
Hypoaspis praesternalis	0	0	0	2.23	0.3	
Leptogamasus tectygynellus	1.07	2.74	2.76	0.74	3.6	
Paragamasus conus	12.3	32.18	38.98	21.34	40.54	
Paragamasus sp.	0	0	0	0.5	0	
Paragamasus vagabundus	10.7	4.02	5.91	0.99	0.6	
Parasitidae sp.	0.53	0	0.39	0.25	0	
Parazercon radiatus	0.53	0.73	0	0	0	
Pergamasus crassipes	0	0.73	0.79	1.49	0	
Pergamasus sp.	0	0	0	0.25	0	
Rhodacarus mandibularosimilis	1.07	0	3.15	1.74	2.1	
Trachytes aegrota	0	0.18	0	0.25	0	
Trichouropoda ovalis	0	1.28	0.79	0	0	
Veigaia cervus	0.53	0	0	0	0.3	
Veigaia nemorensis	2.67	0.91	0.79	0.25	0.9	
Zercon peltatus	1.07	0	0	0	0	
Zercon zelawaiensis	0	1.1	5.91	26.3	7.81	

Table 2. Dominance index (%) of mesostigmatid mites found in transverse belts with different wood residue utilization methods (1 = raked and removed; 2 = burnt; 3 = left undisturbed; 4 = chipped and left on the surface; 5 = chipped and mixed with the soil)

(belt 4). Dominants are represented by *Asca aphidiodes* (belts 4 and 5), *Asca bicornis* (Canestrini & Fanzago, 1887) (belt 2), *Amblyseius* sp. (belts 1, 4 and 5), *Paragamasus vagabundus* (belt 3) and *Zercon zelawaiensis* (belts 3 and 5).

Very high similarity rates (74.58 and 83.42) of dominance indices of all mesostigmatid mite species in the basic belts indicate a low variation in community structure (Fig. 1). It is proved by statistical analysis: the average number of species (Table 3) and the average number of individuals (Table 4) do not differ significantly between



Fig. 1. Similarity dendrogram based on dominance indices of all mesostigmatid mites found in differently prepared basic belts (A = furrow with a ditch; B = no preparation; C = classic furrow)

Table 3. The impact of soil preparation method on the number of species per sample (A = furrow with a ditch; B = no preparation; C = classic furrow)

Preparation method	Average number of species
A	3.1 a
В	3.0 a
С	3.0 a

Average values marked with the same letters do not differ significantly (P > 0.05).

the experimental variants. The similarity rates of all mesostigmatid mites from the transverse belts (Fig. 2, Table 5) are more variable, between 45.14 and 83.57. The statistical analysis (Table 6) shows significant differences between the plots with respect to the dominance of mesostigmatids. The subplot where the residue was raked and removed differs considerably from the other plots with respect to species domi-

Table 4. The impact of soil preparation method on the number of individuals per sample (A = furrow with a ditch; B = no preparation; C = classic furrow)

Preparation method	Average number of individuals
A	13.2 a
В	10.6 a
С	14.4 a

Average values marked with the same letters do not differ significantly (P > 0.05).



Fig. 2. Dendrogram of similarity rates of all mesostigmatid mites found in transverse belts with various methods of wood residue utilization (1 = raked and removed; 2 = burnt; 3 = left undisturbed; 4 = chipped and left on the surface; 5 = chipped and mixed with the soil)

nance. The highest similarity rate (83.56) concerns the subplot where the residue was chipped and mixed with the soil (3) and the subplot where the residue was left undisturbed (1). The wood residue utilization method also influenced the average number of species (the highest for belts 4 and 2, i.e. for grinding the residue and leaving it on the surface and for burning of the residue) (Table 6) and the average number of individuals per sample (definitely the highest for belt 4, i.e. for grinding the residue and leaving it on the surface) (Table 7).

Table 5. Similarity rates of mesostigmatid communities in transverse belts with various methods o
wood residue utilization (1 = raked and removed; 2 = burnt; 3 = left undisturbed; 4 = chipped and
left on the surface; $5 =$ chipped and mixed with the soil)

Transverse belts	4	5	2	1
3	63.137	83.566	75.504	48.722
4		68.780	57.526	45.135
5			73.187	50.655
2				62.482

Table 6. The impact of wood residue utilization method on the number of species per sample (1 = raked and removed; 2 = burnt; 3 = left undisturbed; 4 = chipped and left on the surface; 5 = chipped and mixed with the soil)

Utilization method	Average number of species
1	3.0 ab
2	3.6 a
3	3.0 ab
4	3.9 a
5	1.7 b

Average values marked with the same letters do not differ significantly (P < 0.01).

Table 7. The impact of wood residue utilization method on the number of individuals per sample (1 = raked and removed; 2 = burnt; 3 = left undisturbed; 4 = chipped and left on the surface; 5 = chipped and mixed with the soil)

Utilization method	Average number of individuals
1	9.4 b
2	14.6 ab
3	12.3 ab
4	20.3 a
5	6.9 b

Average values marked with the same letters do not differ significantly (P < 0.01).

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