



## **A comparative study of the fauna associated with nest mounds of native and introduced populations of the red wood ant *Formica paralugubris***

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(Article begins on next page)

1 **A comparative study of the fauna associated with nest mounds of native and introduced**  
2 **populations of the red wood ant *Formica paralugubris***

3  
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47

48 Highlights

49

50 - Comparison of nest fauna associated with native and introduced populations of *Formica*

51 *paralugubris*

52 - Differences in community compositions among sites, nest fauna more similar than soil fauna

53 - Higher oribatid mite diversity in nests than soils, unlike springtails

54 - Several oribatid species associated with nest habitats, whereas only a few springtails

55 - Weak evidence of multiple relocations of mesofauna species during transplants

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57  
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60

61 Abstract

62

63 In the second half of the twentieth century, many red wood ant populations were transferred from  
64 the Alps to the Apennines as biological control agents. Since the introduction involved the  
65 relocation of entire nest mounds, it is presumable that the associated fauna was also relocated.  
66 While the introduction of these ants has raised several concerns about their ecological impact, there  
67 has been no attempt to investigate the introduction of other nest-associated species. In this study, we  
68 collected samples of soil and nest material from three populations of the red wood ant *Formica*  
69 *paralugubris*, one Alpine native and two imported into the Apennines. We aimed to confirm that  
70 nest mounds are hotspots for soil fauna, detect the occurrence of new myrmecophilous species, and  
71 compare the nest-associated fauna among sites, to test the hypothesis of mass species relocation.  
72 We focused our analyses mainly on two taxa, springtails and oribatid mites, two highly  
73 representative groups of the mesofauna inhabiting nest mounds. The results showed higher richness  
74 and diversity in nests than soil for oribatids but not for springtails. We found 17 myrmecophilous  
75 oribatid species, but only two springtail species. Finally, native and imported sites shared only a few  
76 oribatid and springtail species, suggesting that massive relocation did not occur with nest  
77 transplants or it was likely limited. Additionally, we found some species never before collected in  
78 Italy.

79

80 Keywords: red wood ants, myrmecophily, soil fauna, *Formica paralugubris*, oribatid mites,  
81 springtails, new species

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84  
85

## 1. Introduction

86

87 Red wood ants (RWA) are typical in Central and Northern Europe, where seven species of *Formica*  
88 s.s. or the *F. rufa* group are known [1]. In contrast to other ants, these species are mostly associated  
89 with coniferous trees and require cold climates [2]. In the southern portion of their distribution area,  
90 their presence is limited to higher altitudes [3,4,5,6]. In Italy, the species of *Formica* s.s. are  
91 widespread along the Alpine chain, and only the more thermophilic *Formica pratensis* Retzius,  
92 1783 naturally occurs at more southern locations in the Apennine mountains [7], although its exact  
93 distribution is still unknown. In the mid-twentieth century, nests of several alpine RWA species,  
94 such as *F. paralugubris* Seifert, 1996, *F. polyctena* Förster, 1850, and *F. aquilonia* Yarrow, 1955,  
95 were repeatedly introduced to the Apennines as biocontrol agents for forest insect pests, and in  
96 some cases, viable populations established and started to spread [8,9,10]. Their impact on the  
97 arthropod fauna inhabiting the newly occupied area has been demonstrated [9]. All RWA species  
98 are known to profoundly affect local communities, from plants to vertebrates, due to their predatory  
99 ability, dominant status, and capacity to modify the physical properties of the habitats they colonize  
100 [1,11,12,13]. The large aboveground nest mound is one distinguishing characteristic of these ants.  
101 They are formed to maintain stable humidity and temperature in the nest chambers, and often  
102 contrast with harsh external conditions [14,15,16]. For example, in *F. polyctena* mounds, the  
103 difference between the nest and air temperature during the winter can exceed 20 °C [17]. In cold  
104 habitats, this microclimatic stability may turn the RWA mounds into “warm islands”, which can be  
105 attractive for a wide array of organisms [18,19]. More in general, that ant nests can provide shelter,  
106 food and favorable physical conditions for many litter animals has been established for several ant  
107 species [e.g. 20,21,22,23,24].

108 The invertebrate fauna hosted in RWA nest mounds, assessed in multiple studies, includes a  
109 broad range of taxa: isopods, spiders, mites, nematodes, springtails and a long list of insects, from

110 flies to several beetle species [25,26,27,28]. More than 120 obligate myrmecophilous species have  
111 been found in RWA nest mounds, as well as several other species that occasionally inhabit nests  
112 because of their combined and often favorable moisture, pH and temperature conditions [19].  
113 Myrmecophily can be defined as the partial or complete dependence on ant colonies by non-ant  
114 species [29]. This relationship can be based upon parasitic, commensal or even mutualistic  
115 interactions [30,31]. Thus far, more than ten thousand myrmecophilous arthropods are known, and  
116 they have evolved numerous strategies to inhabit or have access to ant nests [32,33,34], such as  
117 chemical camouflage, morphological mimicry and pheromonal attractors [35,36,37].

118 In this study, we analyze the invertebrate fauna hosted in the nest mounds of *F.*  
119 *paralugubris*, a highly polydomous and polygynous RWA species [38]. Since this species was one  
120 of the most often introduced into the Italian peninsula [10], we focus our study on the comparison  
121 of the nest myrmecophiles from native and introduced populations. Ant translocation involved the  
122 movement of entire nest mounds [8], and this likely translocated all the associated fauna as well.  
123 The main aims of this study are: i) to confirm that nest mounds can be attractive for the soil fauna,  
124 both in the native and imported populations, by comparing assemblages inhabiting nest mounds and  
125 soil; ii) to identify potentially myrmecophilous species not previously described in the literature; iii)  
126 to compare soil and nest fauna across sites and evaluate if this difference supports the hypothesis of  
127 species relocation. We focused on springtails (Hexapoda, Collembola) and oribatid mites (Acari,  
128 Oribatida), which were used in previous studies as common representative taxa of the mesofauna  
129 inhabiting RWA nest mounds [39,40,41,42]. Among microarthropods, springtails and oribatid mites  
130 constitute two of the most species-rich taxa of the soil ecosystem [43]. Although quite distant  
131 phylogenetically, the two groups share several ecological and behavioral features because of their  
132 adaptation to similar ecological niches. Both groups feed on litter and soil micro-organisms and  
133 graze on fungi, and thereby affect the dispersion of saprophytic and mycorrhizal species and control  
134 fungal populations [44,45,46,47]. Their community structure is modulated by several natural  
135 factors, including plant diversity, litter quality and meta-population dynamics. However, these two

136 key groups are also sensitive to anthropogenic factors such as land use, soil tillage, environmental  
137 pollution, physical disturbance and fire [48,49,50,51].

138

## 139 **2. Materials and Methods**

140

### 141 *2.1 Study area and sampling design*

142 The sampling was carried out between June and August 2017. Samples of nest material and soil  
143 were collected from three sites, one in the Alps in the Giovetto di Paline Nature Reserve  
144 (abbreviated as GP, 45° 57'57''N, 10° 7'48''E), and two in the Apennines, in the Abetone forest  
145 (abbreviated as AB, 44° 08'50''N, 10° 40'24'' E) and the Campigna Biogenetic Nature Reserve  
146 (abbreviated as CA, within the Foreste Casentinesi, Monte Falterona e Campigna National Park,  
147 43° 52'00''N, 11° 44'14''E) (Fig. 1). The Alpine site, GP, is the site of origin, where nests later  
148 imported to both Apennine sites were collected [52]. Nests used for sampling in the Campigna  
149 Biogenetic Nature Reserve were imported in 1958 [9, 52], whereas the exact year of transplants is  
150 not available for the Abetone forest. However, the period was probably similar (Groppali, personal  
151 communication). The habitats of the three sites are similar, a mixed forest composed of a dominant  
152 conifer species and beech (*Fagus sylvatica* L. 1753). The dominant species in the Alpine site is the  
153 red fir (*Picea abies* (L.) H. Karst, 1881), while in the Apennine sites, it is the white fir (*Abies alba*  
154 Mill, 1759). The geographic position of all ant nests—including sampled nests—was recorded in  
155 five hectares per site by a GPS locator (Garmin eTrex® 10, accuracy ~ 3 m), and the density per  
156 hectare was evaluated.

157 Five nests of *F. paralugubris* of similar size (height range, 1.3–1.5 m; diameter at the base  
158 of the mound, 2.3–2.6 m) were chosen in each area. Nests were spaced at least 15 meters apart to  
159 ensure independent sampling of the nest fauna. Moreover, we preliminarily checked for the  
160 presence of ant trails connecting the selected nests. From each mound, we collected three 1-dm<sup>3</sup>  
161 samples of nest material (total number of samples = 45, 15 per site) and two 1-dm<sup>3</sup> samples of soil

162 three meters from each nest (as control) utilizing a soil corer (15 cm diameter) to a depth of 50 cm.  
163 Nest material was collected at 40 cm depth from the surface of the mound. Each sample was  
164 separately stored in a plastic bag. All samples were transferred to the laboratory within 6 hours from  
165 the time of collection. The material was gently stirred and homogenized in a plastic basin and then  
166 placed into a Berlese funnel. Samples were left in the funnels for 5 days, following Parisi et al. [53].  
167 The specimens collected were examined using a stereomicroscope. Oribatids and springtails were  
168 identified to the species level, while most of the other organisms were identified to a higher  
169 taxonomic level. All samples were stored in pure ethanol for further analysis.

170

## 171 *2.2 Statistical analyses*

172 Species diversity at the three sites was estimated following Chao et al. [54], using as input the  
173 abundance data for oribatids and presence/absence for springtails. The use of presence/absence for  
174 springtails is motivated by the strongly aggregated distribution of several species of this taxon  
175 [55,56], which may hamper diversity estimates [54]. The method for evaluating diversity is based  
176 on the estimation of Hill numbers,  $qD$ , and yields estimates of total (rarefied and extrapolated)  
177 species richness ( $q = 0$ ), and the exponential values of the Shannon diversity ( $q = 1$ ) and Simpson  
178 diversity ( $q = 2$ ) indices. The 95% confidence intervals were obtained from bootstrapping, based on  
179 4999 replications of the reference sample set. Finally, differences in observed species density  
180 among sites (defined as the number of species captured in each sample) were assessed using mixed-  
181 effect modelling with Poisson error distribution, including the nest as a random variable [57]. A  
182 Tukey post-hoc test was then performed to assess the differences in pairs.

183 Association of a species with either nests (i.e., myrmecophily) or soil was evaluated by  
184 analyzing the probability of occurrence in the respective material. Generalized Linear Mixed  
185 Models (GLMM) were used for this analysis, with the type of habitat (nest or soil) as the main  
186 factor. For springtails, we used binomial distribution (presence/absence data) in the model, whereas  
187 for oribatids, given the high difference in abundance between samples, we log-transformed data and

188 used a Gaussian distribution. For springtails, we analyzed only species that occurred in at least three  
189 samples in each site in which they were present. For oribatids, following Elo et al. [42], we tested  
190 only species that occurred in at least three samples and with at least ten individuals per site. Sites in  
191 which the species did not occur were omitted from the analysis.

192 Compositional differences among sites were investigated using multivariate techniques.  
193 Rare species (those with less than three observations) and empty samples with no specimens were  
194 omitted from the following analyses [58]. The multivariate distances among samples were  
195 computed with the Bray-Curtis dissimilarity index after log transformation of the data, and the  
196 resulting distance matrix was analyzed by non-metric multidimensional scaling (nMDS) according  
197 to Clarke and Warwick [58]. Species composition differences were tested with a permutation-based  
198 non-parametric multivariate analysis of variance (PERMANOVA), using the factor “site” (fixed)  
199 and “nest” (random, nested) to account for non-independence of observations.  $\beta$ -diversity was  
200 computed following Anderson et al. [59], and permuting model residuals were compared to  
201 generate a permutation distribution of F under the null hypothesis of no difference in dispersion  
202 between groups. All analyses were conducted using the R software package (ver.3.6) using the  
203 libraries “vegan”, “iNEXT” and “ecodist”.

204

### 205 **3. Results**

206

207 The estimate of nest density was similar among sites (AB ~ 13 nests/ha, CA ~ 12 nests/ha, GP ~ 12  
208 nests/ha). We collected a total of 32 springtail species (11 in GP, 21 in CA and 16 in AB) belonging  
209 to 21 genera, 122 oribatids (49 in GP, 39 in CA and 51 in AB) belonging to 62 genera, 53  
210 morphospecies of other arthropods, 2 morphospecies of Annelida (Haplotaxida) belonging to the  
211 Enchytraeidae and Lumbricidae families, and one nematode morphospecies. Of the 53  
212 morphospecies of arthropods, 39 were insects, subdivided into Coleoptera (23), Diptera (6),  
213 Hemiptera (4), Lepidoptera (2), Hymenoptera (2), and one each of Thysanoptera and Psocoptera.

214 Other arthropods included four Diplopoda, three Chilopoda, three Arachnida, one Protura, one  
215 Diplura and one Symphyla. A detailed list of the morphospecies collected (Collembola and  
216 Oribatida excluded) and their abundance at each site and type of habitat (nest or soil) is reported in  
217 Supplementary material S1.

218         Except for springtails and oribatids, the most represented group was that of beetles. Overall,  
219 most coleopteran morphospecies were found in nests (20 out of 23, 17 of them exclusively in nests),  
220 whereas only seven out of 23 morphospecies were collected in soil samples (three of them  
221 exclusively in soil). Among Coleoptera, Staphylinidae was the most represented taxon, with ten  
222 total morphospecies collected. Diptera (6 morphospecies) were collected only in the soil samples.  
223 All three spider morphospecies collected were associated with nests only. None of the other groups  
224 with more than one morphospecies was exclusive for either of the two types of samples.

225         We found three oribatid species new to Italy: *Damaeus selgae* Pérez-Íñigo, 1966 and  
226 *Scheloribates tuberculatus* Pérez-Íñigo jr., Herrero and Pérez-Íñigo, 1987 only known from Spain  
227 [60], and *Dissorhina ornata peloponnesiaca* Mahunka, 1974, recorded in Spain [61], Greece [62]  
228 and recently, Poland [63]. Moreover, the myrmecophilous beetle *Monotoma conicicollis* Chevrolat,  
229 1837 (Monotomidae), previously recorded only in the Alps, was recently recorded for the first time  
230 in the Apennines [64]. Several species of insects, previously known only in the Alps, have often  
231 been overlooked, and their populations only recently recorded in the northern Apennines and  
232 surrounding areas [65,66].

233         In total, we found 19 species that preferred ant nests (2 springtails and 17 oribatids), whereas  
234 seven species preferred soil (2 springtails and 5 oribatids) (Table 1). Pooling all sites, no difference  
235 in the number of springtail species in soil and nests was found (Fisher exact test  $p = 0.805$ ), whereas  
236 oribatid species richness was significantly higher in nests than soil (Fisher exact test  $p = 0.022$ ). The  
237 number of species shared between soil and nests differed among sites for springtails (Fisher exact  
238 test  $p = 0.022$ , Figure 2), but not for oribatids (Fisher exact test  $p = 0.310$ ). The number of species  
239 shared among sites is summarized in Table 2. There was no difference between the proportion of

240 shared species both in soils and in nests with respect to the total number of species in that habitat  
241 (Mantel-Haenszel chi-square test with continuity correction:  $\chi^2 = 2.1$ ,  $p = 0.147$ ).

242 Focusing on nest samples, CA generally had slightly higher springtail species richness and  
243 diversity than either AB or GP, although the confidence intervals of the three sites overlapped  
244 widely (Figure 3). Species density did not differ among sites for springtails (Type III Wald test:  $\chi^2 =$   
245  $1.727$ ,  $df = 2$ ,  $p = 0.422$ ). For oribatids, CA had lower  $\alpha$ -diversity than AB and GP, while species  
246 density was significantly lower than at AB and GP ( $\chi^2 = 18.943$ ,  $df = 2$ ,  $p < 0.001$ ; multiple  
247 comparisons: CA vs AB:  $z = -3.572$ ,  $p < 0.001$ ; CA vs GP:  $z = -4.078$ ,  $p < 0.001$ ; AB vs GP:  $z = -$   
248  $0.525$ ,  $p = 0.859$ ).

249 The nMDS ordination plot for springtails (Figure 4a, stress = 0.06) showed a separation  
250 among groups, whereas nest samples were partially separated by site. The PERMANOVA (Table 3)  
251 revealed a significant effect of the factor site, but also significant variability among replicate nests  
252 within each site. A clearer picture emerged from the nMDS ordination plot of oribatids (Fig. 4b,  
253 stress = 0.17). In this case, the three sites clearly had different soil assemblages, but the two  
254 introduction sites AB and CA were closer to each other (i.e., more similar) than to GP. The data  
255 points of the former two were located on opposite sides of the plot of the latter one. Nest  
256 assemblages were also fairly distinct by site but were more similar than soil samples. All soil  
257 samples were located in the central part of the plot, in close connection to each other. Notably,  
258 while the soil samples from the Apennine sites were clearly separated by those from the Alps, the  
259 nest samples were more similar. Even in this case, PERMANOVA revealed significant differences  
260 in nest assemblages among sites, both also significant within-site variability (Table 3). Finally, no  
261 significant difference in  $\beta$ -diversity among sites was found for either springtails ( $p = 0.054$ ) or  
262 oribatids ( $p = 0.431$ ).

263

#### 264 4. Discussion

265

266 4.1 Richness and diversity of soil and nest fauna

267 The predictions that higher species richness would be found in nest mounds of *F. paralugubris* than  
268 in the surrounding soil was confirmed for oribatids, but not for springtails. As for this latter group,  
269 the presence of red wood ants is known not to be a determining factor affecting their occurrence  
270 (see Lenoir et al. [40] for a study with *F. polycтена*). However, little is known about the difference  
271 in richness and diversity between ant nests and the surrounding soil. Conversely, for oribatids, our  
272 finding is consistent with Laakso and Setälä [18], who found higher mite richness in nests of *F.*  
273 *aquilonia* than in the nearby soil.

274 Our results contrast with those of Elo et al. [42], who found different species but comparable  
275 species richness in the nest mounds of *F. polycтена* and the surrounding soil. As no other replicate  
276 studies on the same ant species are available, it is not possible to know if these differences represent  
277 specific features of the three species assessed, or, instead, they are driven by local environmental  
278 determinants, as suggested by Elo et al. [42]. These authors hypothesized that this incongruence  
279 could be due to differences in the quality of the leaf litter, which is assumed to be poorer in  
280 coniferous stands than in broadleaf forests. This relative lack of litter might compel mites to  
281 aggregate within nest mounds more strongly in the former than in the latter. A second suggested  
282 factor could be nest density: the higher density may drive species to be more distributed and less  
283 concentrated among ant nests. In our study, the hypothesis regarding leaf litter quality might be  
284 supported, because all three sites were dominated by coniferous trees (*Abies alba* and *Picea abies*),  
285 as in Laakso and Setälä [18], although a denser understory is present in the Alpine site. Nest density  
286 may have an effect as well, but we do not have any reference to verify whether the density in our  
287 sites is relatively high or low, and therefore how it might influence the aggregative behavior of nest  
288 symbionts. However, density was very similar among sites; hence this factor should similarly  
289 influence all three populations.

290

291 4.2 Myrmecophilous or soil-preferring species

292 Springtails and oribatid mites showed different degrees of potential myrmecophily. Only two  
293 springtail species, *Cyphoderus albinus* Nicolet, 1842 and *Lepidocyrtus cyaneus* Tullberg, 1871  
294 selectively chose ant nests, and both are known myrmecophilous inhabitants of RWA nest mounds  
295 [19,26], and other ant species (e.g., *C. albinus* in *Lasius neglectus* Van Loon, Boomsma &  
296 Andrasfalvy, 1990 nests [67]). In contrast, two species, *Folsomia penicula* Bagnall, 1939 and  
297 *Pseudosinella apuanica* Dallai, 1970, both known to occur in Italy [68], preferred soil habitats, even  
298 though in both cases, the preference was not absolute, as they also occurred in some ant nest  
299 samples, though to a lesser extent.

300         Conversely, a higher number of oribatid species (17 out of 122) were found to be potentially  
301 myrmecophilous. Four of these, *Carabodes labyrinthicus* Michael, 1879, *Carabodes ornatus*  
302 Storkán, 1925, *Liacarus coracinus* Koch, 1841, and *Pergalumna nervosa* Berlese 1914, have  
303 previously been described as preferring the nest habitat [42]. Most of the species collected are quite  
304 common, such as the ones belonging to the *Carabodes* genus, which can be found both in European  
305 and American forests, mainly in tree bark where lichens, mosses, and fungi are present [69,70].  
306 Since all these latter organisms (or parts of them) can be found within nest mounds, either alive or  
307 as nesting material [71,72], it is possible that nest habitat is suitable for some *Carabodes* species.  
308 Two *Pergalumna* species were recognized as potentially myrmecophilous, *P. nevosa* and *P. altera*.  
309 Both species have previously been collected in central Italy [73,74]. The species within this genus  
310 are known predators of nematodes [75,76], which, in turn, may find the particular nest conditions  
311 favorable for proliferation [77]. Unfortunately, the Berlese funnel is not an accurate tool for  
312 detecting nematodes, so we cannot provide reliable information on nematode communities.

313         Regarding the linkage between *Phauloppia lucorum* Koch 1841 and ant nests, usually, most  
314 species of the *Phauloppia* genus are known to inhabit and feed on lichens [78,79,80]. Thunes et al.  
315 [81] found that the occurrence of the RWA *F. aquilonia* negatively impacted the lichen community,  
316 probably because of the chemical changes in the environment or by the direct collection of thallus  
317 fragments, which can be found within RWA nest mounds [72]. However, to our knowledge, it is

318 hitherto unclear whether those lichen fragments came from active harvesting by ants or merely from  
319 the collection of detached lichen fragments. In any case, *P. lucorum* may feed on the lichens  
320 transported by ants into the nests, making the nest mound a suitable habitat for this species as well.

321 Only a few species, such as two *Oppiella* species and *Minunthozetes pseudofusiger*  
322 Schweizer, 1922 showed a preference for soils. For the latter, the preference for soil instead of other  
323 habitats, such as grass, roots or litter, is known [82], although studies on their avoidance of ant nests  
324 are not available. The generalist and eurytopic *Oppiella* species are known to occur in harsh  
325 habitats, such as newly burned soils [83], suggesting that in this context, they prefer to occupy  
326 habitats with more extreme temperatures and moisture conditions [84].

327

#### 328 4.3 Comparison of assemblages among sites and species relocation hypothesis

329 We detected differences in species richness and diversity across sites only for oribatids but not for  
330 springtails. Lower oribatid diversity was found at Campigna, whereas the native site of Giovetto di  
331 Paline and Abetone were very similar. This result is quite surprising, because the two imported  
332 populations, Campigna and Abetone, are geographically close and very similar in their general  
333 features (fir-dominated forests, altitude and climate). The Alpine site, although similar in dominant  
334 tree composition, shows denser understory (F. Frizzi, personal observation), which may potentially  
335 affect both springtails and mite diversity [85,86]. However, many factors may influence diversity,  
336 such as altitude, climate conditions, and geographic location [2]. Moreover, metapopulation rules  
337 apply to the nest mound fauna of RWA populations, and inter-nest distances may affect diversity  
338 [87]. Although the mean nest density was similar among sites, the distances between sampled nests  
339 with neighboring ones were not measured in a precise manner, and a different exchange of  
340 mesofauna between nests among sites cannot be ruled out. Thus, the reasons behind this apparent  
341 incongruence are not yet determined and could be a subject for future studies.

342 According to the multivariate analysis, the three sites had different springtail and oribatid  
343 assemblages, both for soil and nest communities. No springtail species and only three oribatid

344 species were shared among the three soil sites. The difference in the soil assemblages probably  
345 reflects the geographic distance, which is shorter between the two imported populations than  
346 between native and imported sites. On the other hand, notably, the oribatid fauna was more similar  
347 among nests than among soils, irrespective of the site, which may suggest a wider geographic  
348 diffusion of some nest-preferring species than of the soil fauna. Indeed, the proportion of shared  
349 species was nearly twice among nest than soil samples (12 out of 102 in nests, 3 out of 54 in the  
350 soil). To our knowledge, this is a novel finding and a targeted genetic analysis of these taxa in soils  
351 and nest mounds could shed light on the effective dispersion history of the species [88,89].

352           Nonetheless, the number of species sampled in nests and shared among sites (2 springtails  
353 and 12 oribatids) was rather low compared to the total number of species collected in all nest  
354 samples (22 springtails and 102 oribatids). This result may suggest that the number of species  
355 transferred from the Alps and established in the Apennines was not high or that some of the species  
356 already occurred in central Italy. For example, the two springtail species, *C. albinus* and *L. cyaneus*,  
357 are known myrmecophilous species in many parts of Europe [26,67]. Hence it is likely that they  
358 spontaneously occur at all sites independent of introductions. Thus, possible relocation with nest  
359 material can be hypothesized only for a few species. For example, we collected a few specimens of  
360 the oribatid mite *Jugatala angulata* Koch, 1839 in the nests of all three sites, even though this  
361 species has never been recorded in Italy except for the Alps [90,91]. Similarly, *Cymbaeremaeus*  
362 *cymba* Nicolet, 1855 was found only in nest samples in both introduction sites, although this species  
363 has already been recorded in the Italian peninsula, but only in northern and southern sites and not in  
364 the central regions [73]. For the other nest-preferring species, it is challenging to hypothesize a  
365 relocation process, because some of them were also recorded in soil samples (*Adoristes ovatus*  
366 Koch, 1839, *C. labyrinthicus*, *Hermannia gibba* Koch, 1839, *Scheloribates pallidulus* Koch, 1841).  
367 It cannot be excluded that individuals of some species—not strictly dependent on ants to survive—  
368 shifted toward inhabiting the soil.

369

370       **5. Conclusions**

371

372   In conclusion, this is the first study to include an introduced population in a comparative analysis of  
373   the nest-associate fauna in red wood ants. We confirmed that RWA nests are hotspots for arthropod  
374   biodiversity, particularly for oribatid mites. We found several potentially myrmecophilous species  
375   that preferred the nest habitat instead of soil, though the symbiotic relationships with ants should be  
376   further investigated to verify myrmecophily. We used for the first time a comparative approach in  
377   the analysis of the nest fauna, assessing the differences between a native Alpine population and two  
378   introduced Apennine populations of the RWA *F. paralugubris*. Although we might expect some  
379   clear traces of a large species relocation given the massive transport of nest material during  
380   introductions, we found only a few species that potentially could have been transferred, suggesting  
381   that it probably did not occur. More comparative studies are recommended, e.g. on RWA imported  
382   populations, since the fate of most of them—and the fauna inhabiting their nests—are hitherto  
383   almost wholly unknown.

384

385 **Fig. 1.** Map of the three sampling sites. In light grey is the alpine site of the original population, and  
386 in dark grey, the two Apennine sites of the two introduced populations.

387

388 **Fig. 2.** Cumulative number of species observed at the three study sites for a) springtails and b)  
389 oribatids. Black bars = nests, white bars = soil, grey bars = species shared between soils and nests.  
390 Sites: AB = Abetone (Apennines), CA = Campigna (Apennines), GP = Giovetto di Paline (Alps).

391

392 **Fig. 3.** Species diversity in the three sites (red = Abetone, Apennines; blue = Campigna, Apennines;  
393 green = Giovetto di Paline, Alps). Sample-based plots with 95% confidence intervals for species  
394 richness ( $q=0$ ), Shannon ( $q=1$ ), and Simpson ( $q=2$ ) diversity indexes. Continuous lines = observed  
395 values; dashed lines = extrapolated values.

396

397 **Fig. 4.** nMDS ordination plots for a) springtail and b) oribatid datasets. Colors indicate the sites (red  
398 = Abetone, Apennines; blue = Campigna, Apennines; green = Giovetto di Paline, Alps), symbols  
399 indicate sample types (triangles = samples from nests; circles = samples from soils).

400

401  
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403

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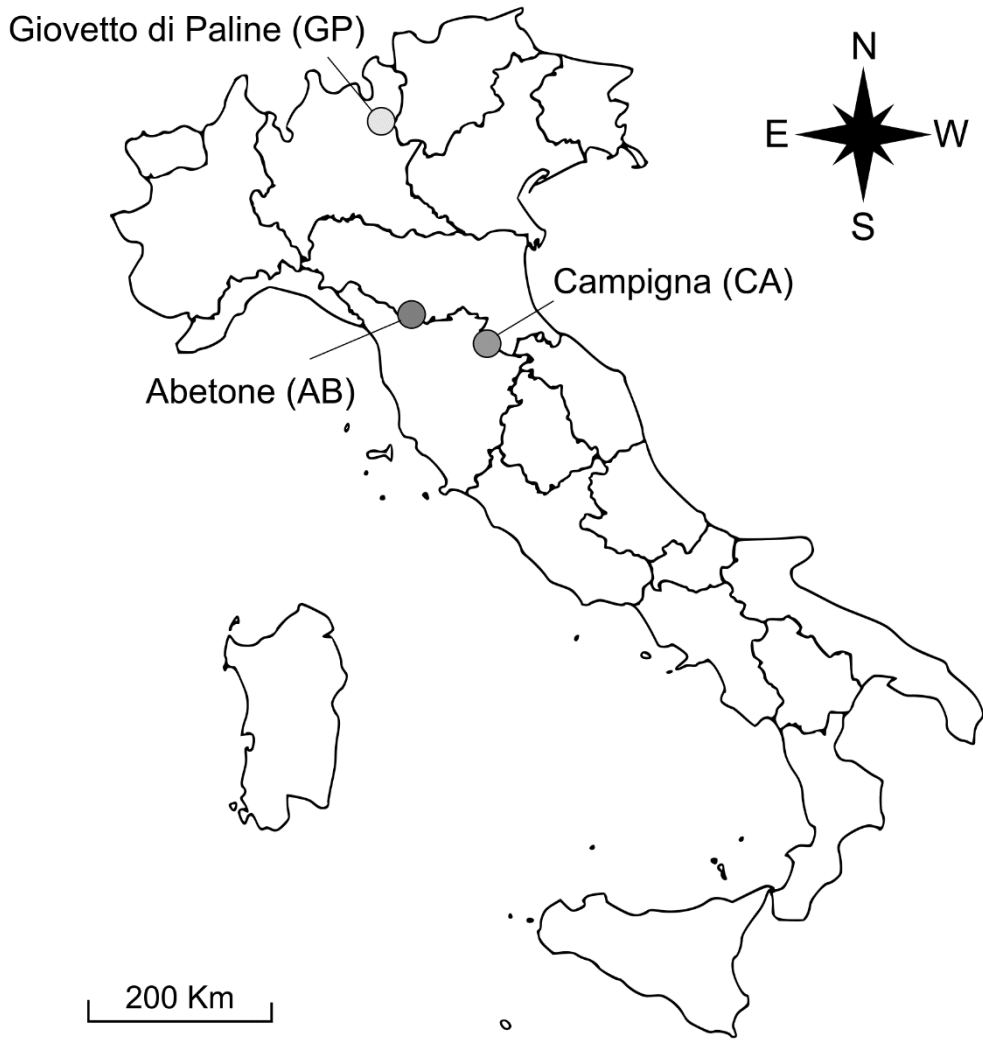
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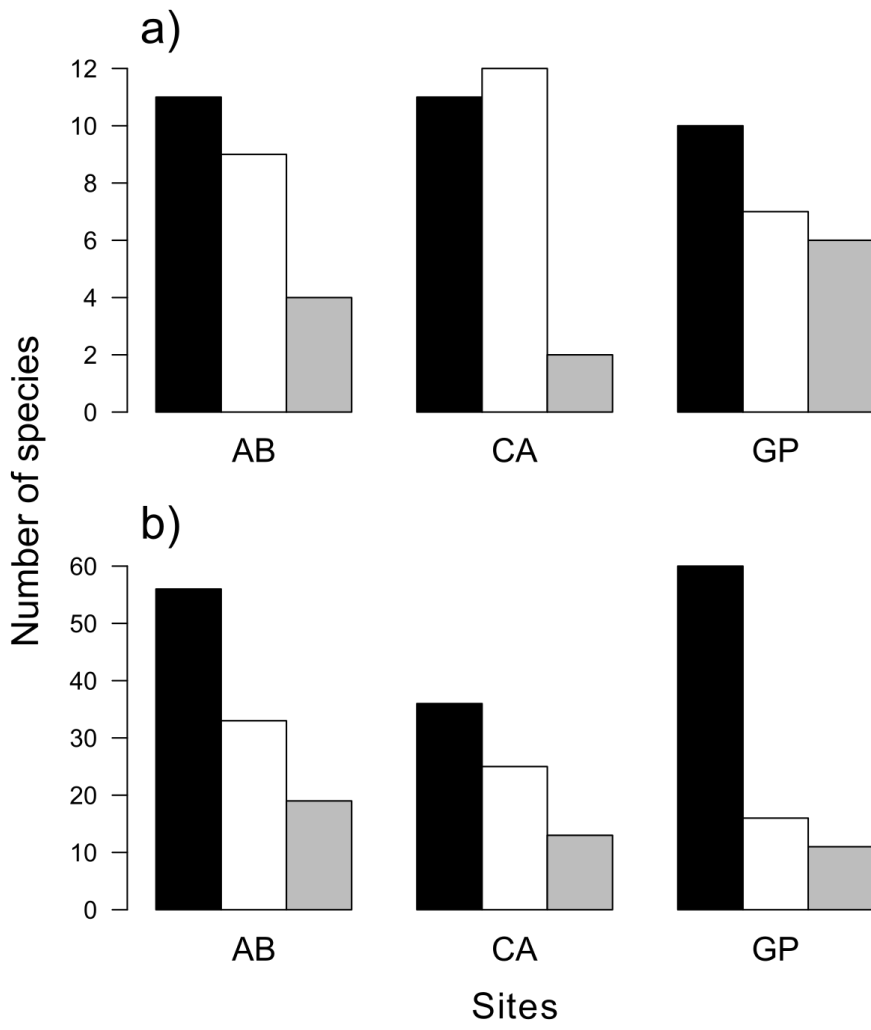
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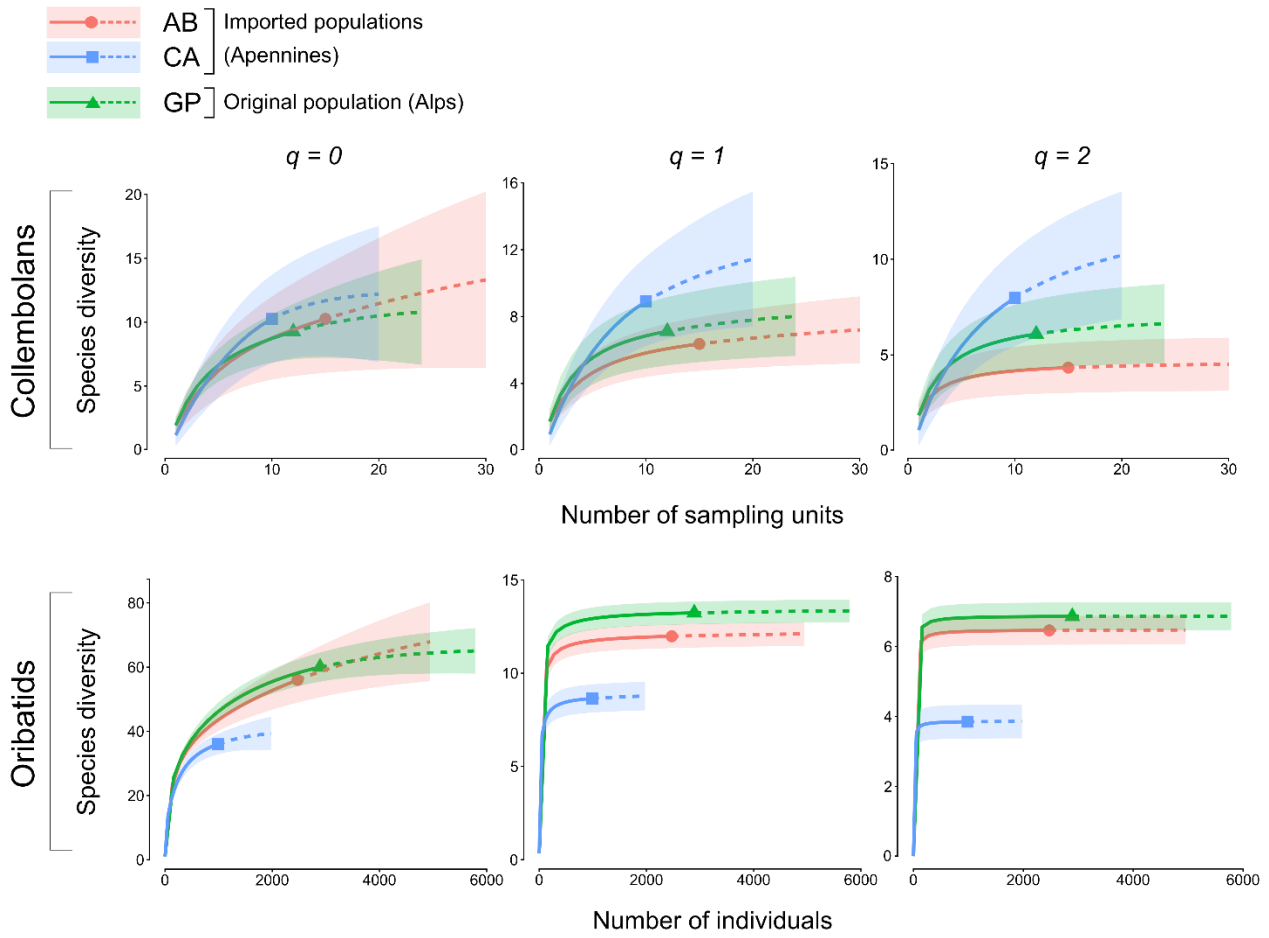
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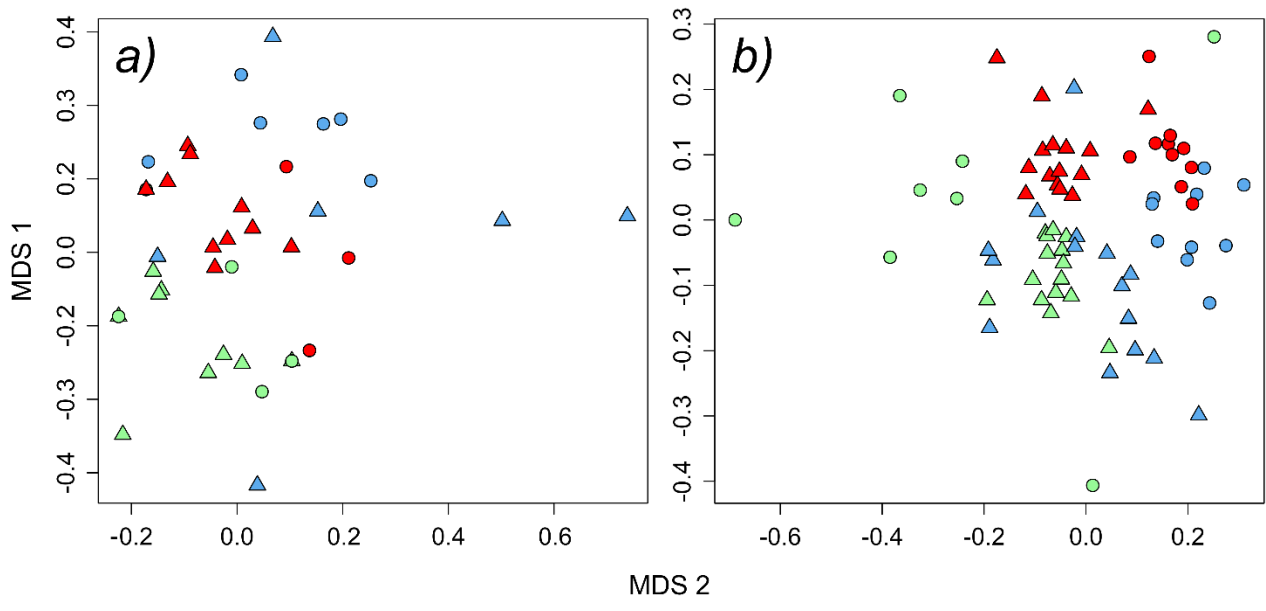




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669 **Table 1**

670 Species of springtails and oribatids collected, with their abundance and the estimated preference for

671 nest or soil according to results of models. Pref = preference. Sites: AB = Abetone (Apennines), CA

672 = Campigna (Apennines), GP = Giovetto di Paline (Alps).

673

Springtails Species	AB		CA		GP		Pref
	Nest	Soil	Nest	Soil	Nest	Soil	
<i>Anurophorus atlanticus</i>	1	-	-	-	-	-	
<i>Ceratophysella armata</i>	-	2	-	10	-	-	
<i>Cyphoderus albinus</i>	2277	-	24	-	438	2	Nest
<i>Desoria</i> sp.	1	-	9	1	-	-	
<i>Entomobrya nivalis</i>	-	-	-	-	4	-	
<i>Entomobrya</i> sp.	3	2	1	-	-	-	
<i>Folsomia manolachei</i>	6	-	-	12	-	-	
<i>Folsomia penicula</i>	5	39	-	-	32	25	Soil
<i>Folsomia</i> sp.	-	-	-	3	-	-	
<i>Friesea mirabilis</i>	24	-	6	-	-	-	
<i>Hypogastrura viatica</i>	-	-	-	-	1	-	
<i>Isotomurus</i> sp.	-	-	-	3	-	-	
<i>Lathriopyga longiseta</i>	-	-	3	-	-	-	
<i>Lepidocyrtus violaceus</i>	-	-	9	-	-	-	
<i>Lepidocyrtus cyaneus</i>	8	-	21	-	56	1	Nest
<i>Lepidocyrtus lanuginosus</i>	-	-	-	-	1	-	
<i>Lepidocyrtus lignorum</i>	-	-	-	-	1	1	
<i>Mesaphorura</i> sp.	-	-	-	-	-	1	
<i>Paratullbergia callipygos</i>	-	1	1	-	-	-	

<i>Parisotoma notabilis</i>	18	1	-	-	6	1	
<i>Protaphorura armata</i>	-	1	1	1	-	-	
<i>Protaphorura campata</i>	-	1	-	-	-	-	
<i>Protaphorura cancellata</i>	-	-	2	-	6	1	
<i>Pseudachorutes dubius</i>	-	-	-	2	-	-	
<i>Pseudachorutes sp.</i>	-	-	-	1	-	-	
<i>Pseudosinella alba</i>	-	-	-	1	-	-	
<i>Pseudosinella apuanica</i>	6	13	-	1	-	-	Soil
<i>Tetracanthella pilosa</i>	-	-	-	2	-	-	
<i>Tomocerus minor</i>	-	2	-	-	-	-	
<i>Vertagopus arborea</i>	-	-	2	-	-	-	
<i>Xenylla maritima</i>	-	-	-	-	4	-	
<i>Xenylla schillei</i>	29	-	-	1	-	-	

Oribatids	AB		CA		GP		Pref
	Nest	Soil	Nest	Soil	Nest	Soil	
<i>Achipteria coleoptrata</i>	-	-	25	7	19	-	
<i>Achipteria italica</i>	-	-	6	2	-	-	
<i>Acrogalumna longipluma</i>	55	-	32	-	-	-	Nest
<i>Adoristes ovatus</i>	19	-	64	2	111	1	Nest
<i>Atropacarus clavigerus</i>	-	-	-	-	19	-	
<i>Atropacarus platakisi</i>	-	-	-	-	3	-	
<i>Banksinoma lanceolata</i>	-	-	1	-	-	-	
<i>Belba bartosi</i>	-	-	-	-	1	-	
<i>Berniniella (Berniniella) hauseri</i>	3	-	-	-	-	-	
<i>Berniniella bicarinata</i>	7	1	-	14	-	-	
<i>Berniniella coronata</i>	-	-	-	-	-	1	
<i>Berniniella (Hypogeoppia) dungeri</i>	-	9	-	-	-	-	

<i>Camisia horrida</i>	-	-	-	-	2	-	
<i>Camisia spinifer</i>	4	-	-	-	-	-	
<i>Carabodes areolatus</i>	2	-	-	-	2	-	
<i>Carabodes labyrinthicus</i>	149	3	3	5	13	1	Nest
<i>Carabodes manganoi</i>	3	-	4	-	-	-	
<i>Carabodes marginatus</i>	-	-	4	-	5	-	
<i>Carabodes ornatus</i>	11	-	5	-	19	-	Nest
<i>Carabodes palmifer</i>	1	-	-	-	-	-	
<i>Carabodes reticulatus</i>	-	-	-	-	1	-	
<i>Cepheus pegazonae</i>	1	-	-	-	-	-	
<i>Cerachipteria minuscula</i>	23	1	1	8	-	-	
<i>Ceratozetes gracilis</i>	-	-	59	21	2	-	
<i>Ceratozetes minutissimus</i>	-	-	-	-	2	-	
<i>Ceratozetes peritus</i>	2	13	-	-	-	-	Soil
<i>Chamobates cuspidatus</i>	-	1	-	-	1	-	
<i>Chamobates pusillus</i>	23	1	-	-	-	-	
<i>Chamobates rastratus</i>	-	-	-	-	271	-	
<i>Chamobates voigsti</i>	-	-	-	-	5	6	
<i>Cymbaeremaeus cymba</i>	9	-	3	-	1	3	
<i>Damaeus selgae</i>	-	-	-	-	60	-	
<i>Dissorhina ornata peloponnesiaca</i>	-	4	-	-	-	-	
<i>Eniochthonius minutissimus</i>	-	-	-	-	24	-	
<i>Eremaeus cordiformis</i>	-	-	4	-	5	-	
<i>Eueremaeus oblongus</i>	340	-	-	-	-	-	Nest
<i>Eueremaeus silvestris</i>	-	-	52	9	-	-	
<i>Eueremaeus valkanovi</i>	-	-	-	-	7	-	
<i>Eulohmannia ribagai</i>	-	-	-	1	-	-	

<i>Eupelops acromios</i>	3	-	-	-	3	-	
<i>Eupelops affinis</i>	1	-	-	-	-	-	
<i>Eupelops torulosus</i>	1	-	1	-	46	-	Nest
<i>Euzetes globulus</i>	-	1	-	-	-	-	
<i>Fosseremus laciniatus</i>	-	-	-	-	-	1	
<i>Hemileius initialis</i>	-	-	-	-	1	-	
<i>Hermannia gibba</i>	48	1	2	2	29	-	
<i>Hypochthonius luteus</i>	-	-	-	-	8	-	
<i>Jugatala angulata</i>	2	-	2	-	1	-	
<i>Lepidozetes singularis</i>	-	-	6	-	6	2	
<i>Liacarus coracinus</i>	30	1	-	-	197	3	Nest
<i>Liacarus perezinigo</i>	-	-	-	-	1	-	
<i>Liacarus subterraneus</i>	-	-	4	-	-	-	
<i>Liebstadia humerata</i>	-	-	-	-	3	-	
<i>Liebstadia longior</i>	15	-	-	-	-	-	Nest
<i>Medioppia pinsapi</i>	-	-	-	-	-	2	
<i>Metabelba parapulverosa</i>	-	-	-	-	35	-	
<i>Micropia minus</i>	-	7	-	-	-	-	
<i>Minunthozetes pseudofusiger</i>	5	75	-	53	-	-	Soil
<i>Multioppia glabra</i>	-	-	-	-	5	-	
<i>Nanhermannia nana</i>	-	-	-	-	1	-	
<i>Neotrichoppia (Confinoppia) confinis</i>	20	9	-	3	84	1	
<i>Nothrus anauniensis</i>	-	1	-	1	-	-	
<i>Nothrus silvestris</i>	-	-	-	-	-	1	
<i>Odontocephus elongatus</i>	11	-	-	-	2	-	
<i>Ommatocephus ocellatus</i>	5	-	-	-	-	-	
<i>Ophidiotrichus vindobonensis</i>	-	-	-	-	3	-	

<i>Oppiella (Moritzoppia) keilbachi</i>	-	-	-	-	140	-	
<i>Oppiella (Moritzoppia) uncarinata</i>	-	-	37	-	4	-	
<i>Oppiella (Oppiella) acuminata</i>	96	-	-	-	-	-	
<i>Oppiella (Oppiella) nova</i>	30	39	-	10	25	-	
<i>Oppiella (Rhinippia) fallax</i>	1	2	-	-	-	-	
<i>Oppiella (Rhinoppia) obsoleta</i>	-	39	-	-	-	-	Soil
<i>Oppiella (Rhinoppia) similifallax</i>	-	1	-	-	-	-	
<i>Oppiella (Rhinoppia) subpectinata</i>	1	97	-	95	8	-	Soil
<i>Oribatella brevipila</i>	-	-	11	-	58	-	
<i>Oribatella calcarata</i>	-	-	-	-	4	-	
<i>Oribatella euthricha</i>	-	-	2	-	-	-	
<i>Oribatella hungarica</i>	15	-	-	-	-	-	
<i>Oribatella quadricornuta</i>	-	-	-	-	7	-	
<i>Oribatella sexdentata</i>	1	-	-	-	-	-	
<i>Oribatula tibialis</i>	4	-	8	3	2	1	
<i>Pantelozetes paolii</i>	-	1	-	-	2	-	
<i>Parachipteria petiti</i>	13	11	-	-	-	-	
<i>Parachipteria punctata</i>	1	-	-	-	-	-	
<i>Pergalumna altera</i>	39	-	4	-	63	-	Nest
<i>Pergalumna nervosa</i>	131	-	83	-	736	-	Nest
<i>Phauloppia lucorum</i>	17	-	-	-	-	-	Nest
<i>Phauloppia rauschenensis</i>	-	-	-	-	2	-	
<i>Phthiracarus anonymus</i>	1	-	-	-	8	-	
<i>Phthiracarus clavatus</i>	2	-	1	-	-	-	
<i>Phthiracarus globosus</i>	-	-	1	-	-	-	
<i>Phthiracarus italicus</i>	-	1	-	3	-	-	
<i>Phthiracarus laevigatus</i>	-	-	9	-	-	-	

<i>Phthiracarus lentulus</i>	1	1	-	-	-	-	
<i>Phthiracarus nitens</i>	1	2	5	1	-	-	
<i>Phthiracarus peristomaticus</i>	-	-	-	-	12	-	
<i>Platynothrus peltifer</i>	-	-	-	-	1	-	
<i>Porobelba spinosa</i>	551	-	121	-	-	-	Nest
<i>Poroliodes farinosus</i>	2	-	-	-	-	-	
<i>Ramusella (Inscuptoppia) elliptica</i>	1	22	-	-	-	-	Soil
<i>Ramusella (Ramusella) assimilis</i>	-	-	3	4	1	-	
<i>Raphigneta numidiana</i>	-	-	-	-	-	1	
<i>Rhysotritia duplicata</i>	-	-	10	1	109	1	Nest
<i>Scheloribates laevigatus</i>	1	-	-	-	-	-	
<i>Scheloribates pallidulus</i>	565	1	407	-	574	-	Nest
<i>Scheloribates tuberculatus</i>	-	-	-	-	108	3	
<i>Serratoppia serrata</i>	-	1	-	-	-	-	
<i>Steganacarus (Steganacarus) applicatus</i>	-	-	-	-	25	-	
<i>Steganacarus (Steganacarus) donatoi</i>	-	1	-	-	-	-	
<i>Steganacarus (Steganacarus) hirsutus</i>	19	1	-	-	-	-	Nest
<i>Steganacarus (Tropacarus) carinatus</i>	1	-	-	-	2	-	
<i>Suctobelbella acutidens</i>	-	1	-	-	-	-	
<i>Suctobelbella alleonasuta</i>	-	-	-	2	-	-	
<i>Suctobelbella cf. subcornigera</i>	1	-	-	-	-	-	
<i>Suctobelbella opistodentata</i>	-	-	-	2	-	-	
<i>Suctobelbella sarekensis</i>	-	1	-	2	-	-	
<i>Tectocepheus sarekensis</i>	9	1	-	5	1	1	
<i>Tectocepheus velatus</i>	10	-	2	1	2	-	
<i>Xenillus clypeator</i>	-	-	1	-	-	-	
<i>Xenillus tegeocranus</i>	22	-	-	-	-	-	Nest

<i>Zygoribatula exilis</i>	143	-	4	-	-	-	Nest
<i>Zygoribatula propinqua</i>	1	-	-	-	-	-	

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677 **Table 2**

678 Number of species shared between sites, both for soils and nests. Sites: AB = Abetone (Apennines),

679 CA = Campigna (Apennines), GP = Giovetto di Paline (Alps).

680

Group	Springtails				Oribatids			
	soil		nests		soil		nests	
	shared	total	shared	total	shared	total	shared	total
AB-CA	3	18	5	17	13	45	19	73
AB-GP	2	14	4	17	4	45	22	94
CA-GP	0	19	3	18	6	35	21	75
AB-CA-GP	0	23	2	22	3	54	12	102

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683 **Table 3**

684 Results of PERMANOVA tests.

Factor	Springtails				Oribatids			
	df	SS	F	<i>P</i>	df	SS	F	<i>P</i>
Site	2	2.300	2.35	< 0.001***	2	3.669	4.04	< 0.0001***
Nest[Site]	12	5.864	3.12	< 0.0001***	12	5.451	2.24	< 0.0001***
Residuals	21	3.288			30	6.076		

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