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## Functional role of microarthropods in soil aggregation

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## 8 **Functional role of microarthropods in soil aggregation**

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26 **Soil aggregation has received a lot of attention in the last years; however, the**  
27 **focus was mostly on soil microorganisms or larger soil fauna, especially**  
28 **earthworms. The impact of the large group of microarthropods, e.g. Collembola**  
29 **and Acari, is nearly unknown and hence underrepresented in the literature. Here**  
30 **we propose and discuss potential direct and indirect mechanisms of how**  
31 **microarthropods could influence this process with the focus on collembolans,**  
32 **which are in general a relatively well studied taxon. Indirect mechanisms are**  
33 **likely to have larger impacts on soil aggregation than direct effects. The variety**  
34 **of indirect mechanisms based on the provision of organic material like faecal**  
35 **pellets, molts and necromass as food source for microorganisms is high and**  
36 **given available evidence we propose that these mechanisms are the most**  
37 **influential. We highlight the need for overcoming the challenges of culturing and**  
38 **handling of these animals in order to be able to design small scale experiments**  
39 **and field studies which would enable us to understand the role of the different**  
40 **functional groups, their interaction with other soil fauna and the impact of land**  
41 **use practices on soil aggregation.**

42 *Key words: soil structure; microarthropods; Collembola; Acari*

43

#### 44 ***Introduction***

45 Soil structure plays a critical ecosystemic role in biogeochemical processes (e.g.  
46 Jastrow, 1996), water infiltration, gas exchange efficacy, and resistance against  
47 erosional loss, and influences the performance of soil biota, including roots (Hartge and  
48 Stewart, 1995; Miller and Jastrow, 1992; Oades, 1984; Rillig and Mummey, 2006). Soil  
49 structure is often referred to as the arrangement of different macro- and microaggregate

50 size fractions (organic/mineral complexes of  $>250\mu\text{m}$  or  $<250\mu\text{m}$ , respectively) and the  
51 corresponding pore spaces (Hartge and Stewart, 1995; Rillig and Mummey, 2006).  
52 In hierarchically structured soils, organic matter serves as the main binding agent to  
53 form and stabilize aggregates (Tisdall and Oades, 1982), but additionally, soil texture,  
54 soil microorganisms, roots, inorganic binding agents, the predominant environmental  
55 conditions, and the soil fauna are important for this process (Dexter and Horn, 1988;  
56 Rillig et al., 2015).

57 While soil fauna is generally acknowledged as being important for soil aggregation,  
58 direct empirical evidence is scarce for microarthropods, including mites and  
59 collembolans, the two most abundant and diverse groups. This is surprising given that  
60 these animals can occur at high densities, and given their role in the processing of  
61 organic matter via chemical, physical and biological mechanisms (Lee and Foster,  
62 1991; Wolters, 2000). We are only aware of two studies that have experimentally  
63 quantified the impact of Collembola on soil structure (Siddiky et al., 2012a, b); these  
64 experimental data, however, revealed an effect size comparable to that of much more  
65 thoroughly studied soil biota, such as fungi. These experiments should be extended to  
66 the field as this might also be of agricultural interest.

67 Among the various groups of soil biota, especially the effects of mycorrhizal fungi,  
68 bacteria, earthworms, and termites have been studied intensely (e.g.; Lee and Foster,  
69 1991; Oades and Waters, 1991; Bossuyt et al., 2005; Pulleman et al., 2005; Rillig and  
70 Mummey, 2006; Velasquez et al., 2007). It is known that the excretion of e.g.  
71 polysaccharides by bacteria and the physical enmeshment of soil particles by fungal  
72 mycelia have a positive effect (see e.g. Degens, 1997; Lynch and Bragg, 1985; Oades,  
73 1993; Rillig and Mummey, 2006; Tisdall, 1994b). Larger soil animals like earthworms  
74 and termites directly affect soil structure by their burrowing activities and by the

75 digestion and excretion of relatively large amounts of organic material and soil  
76 particles, which might also lead to increased soil aggregation (e.g. Lavelle, 1988; Lee,  
77 1985; Lee and Foster, 1991; or see review by Six et al., 2004; Tisdall, 1994a, 1994b).  
78 Given this striking asymmetry in our understanding of biotic contributions to soil  
79 aggregation, we here propose and discuss potential mechanisms for Collembola, which  
80 are also likely applicable to other soil microarthropods. We distinguish between direct  
81 and indirect effects (Fig. 1); however all the mechanisms we discuss would in reality  
82 take place simultaneously and in interaction with each other. As the collembolan  
83 *Folsomia candida* is very well studied, especially with regard to properties that might  
84 be involved in mechanisms of soil aggregation, we base our discussion mostly on this  
85 species, but we believe without much loss of generality.

86

87 [Fig. 1]

88

### 89 ***Direct mechanisms***

90 Direct effects of collembolans on soil structure can be categorized in terms of input of  
91 organic material, which positively contributes to soil structure, and degradation of  
92 aggregates, which is a negative effect.

### 93 *Organic matter inputs*

94 Possible positive, direct effects of collembolans on soil structure include the  
95 production, modification and movement of organic matter, which can then serve as  
96 binding agents, nuclei or building blocks for aggregates. Assimilated nutrients can  
97 either be contained in animal tissue or be excreted as metabolic waste. Especially

98 because microarthropods can occur in high numbers, they might produce a large  
99 amount of faecal pellets. It has to be assumed that many soils contain millions of faecal  
100 pellets per square meter (Hopkin, 2007). In this context, Kubiena (1953) reports about  
101 the so-called 'alpine pitch rendzinas' on limestone which are nearly completely  
102 composed of collembolan faeces forming a 15-20cm deep black humus layer.

103 Collembolan eggs are deposited in clutches and need a couple of days to weeks to  
104 develop (Hopkin, 2007). Eggs of the collembolan family Sminthurida might be covered  
105 by a mixture of soil and collembolan waste to protect them from mold and dehydration  
106 (Betsch-Pinot, 1976, 1977; Dallai et al., 2008). After hatching, the remaining egg  
107 integuments might serve as source of fresh organic material to microorganisms (which  
108 will be discussed in the paragraph about indirect mechanisms) or, due to the attached  
109 soil particles and organic material, as nuclei for microaggregate formation.  
110 Collembolans go through several instars, which might mean molting at fairly high rates.  
111 Most species molt throughout their whole life (up to 45 times). Specimens of *Folsomia*  
112 *candida* may live up to six months; however, for other species shorter or far longer  
113 (one year and longer) life-spans have been reported (Hopkin, 2007), which means that  
114 their production of molts could be significant. Interestingly, some oribatid mites can  
115 even survive for up to three years (Capinera, 2008). Their molts are hard-bodied due  
116 to chitin and other components in the cuticle (see Weigmann, 2006) and hence their  
117 breakdown should be slower, and thus they could serve as more long-lived building  
118 blocks of aggregates. Finally, the production of necromasses especially in short-lived  
119 species besides faecal pellets, molts and eggs, can potentially influence soil  
120 aggregation. Unfortunately, there is no study dealing explicitly with the input of these  
121 types of organic material. Given the potentially high local abundances, this should  
122 clearly be a target of future research.

123 *Degradation of soil aggregates via disturbance*

124 Collembola and Oribatida usually consist of populations in the order of 10,000 to  
125 100,000 individuals per square meter (see Hopkin, 2007; Weigmann, 2006). Can they  
126 therefore counteract the formation of aggregates by crawling around or feeding on e.g.  
127 microorganisms, plant remains or various excretory products? The impact of this  
128 disturbance on a *per capita* basis might be low, but data about the impact of locally  
129 high abundant microarthropods on soil aggregation are missing.

130

131 ***Indirect mechanisms***

132 Several studies have investigated the soil food web, functional characteristics and  
133 feedbacks between the different organism groups also in relation to aboveground  
134 biota; however, there are few data on the impact of interacting taxa like fungi and  
135 microarthropods on soil aggregation (Salmon and Ponge, 2001, Siddiky et al., 2012a,  
136 2012b). Fungi and bacteria are directly and indirectly contributing to the production and  
137 release of materials and compounds that contribute to soil structure dynamics while soil  
138 animals affect the translocation and provision of organic material for colonization, like  
139 faecal pellets, molts, eggs, and necromass, and the modification of the activity of  
140 microorganisms by grazing (Coleman et al., 2002). There are studies suggesting that  
141 Collembola could have a positive effect on mycorrhizal functioning as their fungal  
142 grazing might enhance fungal growth and respiration (Lussenhop, 1992). Other studies  
143 suggest that collembolans could also have no or negative effects (Fitter and Garbaye,  
144 1994; Fitter and Sanders, 1992), which brought attention to collembolans as important  
145 regulators of the mycorrhizal symbiosis acting in a density-dependent fashion (Gange,  
146 2000). If there were positive effects on fungal growth or branching patterns, these

147 effects could enhance soil aggregation processes, while the reduction of fungal  
148 biomass could have either negative effects or change the composition of the soil  
149 microbial community with unclear functional consequences. It is also likely that the  
150 observed effects depend on the abundance of Collembola or other microarthropods, a  
151 hypothesis that should therefore be tested (for enchytraeids see Hedlund and  
152 Augustsson, 1995). It has also been shown that Collembola do feed on  
153 arbuscularmycorrhizal fungi (AMF), but, depending on the species, prefer non-AMF  
154 mycelia (e.g. Klironomos and Kendrick, 1996; Klironomos and Ursic, 1998; Moore et  
155 al., 1985; Thimm and Larink, 1995). Another important aspect of the interaction  
156 between the microbial community and microarthropods is the dispersal of spores  
157 (Klironomos and Moutoglis, 1999; Lussenhop, 1992). AMF spores can be far larger  
158 (20-500  $\mu\text{m}$ ) than non-AMF spores (Trappe, 1982) and it is more likely that spores are  
159 ingested by earthworms rather than by Collembola (Fitter and Sanders, 1992; Moore  
160 et al., 1985). Brown (1995) has shown that spores can survive the gut passage of  
161 earthworms with an increased germination rate afterwards (for more information about  
162 gut microbiota in various taxa see e.g. Pherson and Beattie, 1979; Ponge  
163 and Charpentie, 1981, König, 2006.). Still, collembolans are also able to act as vectors  
164 by transporting spores attached to their cuticle (Gormsen et al., 2004), which is also  
165 known for one oribatid group, the Damaeidae (Weigmann, 2006). Although this  
166 phenomenon might be restricted to only a few species it should be considered as  
167 important means of microbial transport which might have an impact on the composition  
168 of the microbial community.

169 As described in the paragraph about the provision of organic material, one major  
170 question is how the organic materials influence the colonization by and composition of



171 microbial communities, which might lead to enhanced aggregate formation. There are  
172 several potential mechanisms which have been investigated only in part so far.

173 Foster et al. (1983) report that fresh faecal pellets can be recognized as round and  
174 smooth surfaces under the scanning electron microscope, whereas older pellets are  
175 mostly densely covered by fungal hyphae; this highlights the importance of  
176 microarthropods in assisting microbial colonization of organic matter. It is also known  
177 that during the molting of collembolans the whole midgut epithelium is also excreted to  
178 dispose of the accumulated toxins (Fountain and Hopkin, 2001; Humbert, 1979). The  
179 total gut volume of *Folsomia candida* was estimated up to 10nl, faecal pellets had a  
180 volume of approx. 1nl (Thimm et al., 1998) and contained approx.  $1.55 \times 10^4$  bacterial  
181 cells (identified by light microscopy) of which only less than 0.01% were dead. Taking  
182 into consideration that, under laboratory conditions, the reported period between the  
183 ingestion and the defecation of bacterial cells can be less than one hour (Czarnetzki  
184 and Tebbe, 2004), the amount of living microbial cells excreted per individuum during  
185 a life cycle is enormous. Some authors (e.g. Hanlon, 1981; Thimm et al., 1998) have  
186 already highlighted the importance of the constant local input of gut (but also other  
187 ingested) bacteria which might lead to an enhanced competition between already  
188 existing soil microorganisms, and this might affect soil aggregation depending on the  
189 ensuing species composition. The same might also be true for other organic material  
190 provided through oviposition or necromass.

191 Collembolans usually excrete urine via the labial nephridia, but can also release  
192 insoluble products via the midgut epithelium (Hopkin, 1997; Larsen, 2007). Most of the  
193 nitrogenous and phosphorus-containing waste products are released as ammonia  
194 (Sjursen and Holmstrup, 2004), uric acid and phosphate, depending on the species. In  
195 spite of the studies addressing these aspects (e.g. Cragg and Bardgett, 2001; Milcu et

196 al., 2006; Verhoef et al., 1988) it is not clear how these different waste  
197 products influence local environmental conditions and hence the microbial  
198 community. Some studies have also investigated partly species-specific characteristics  
199 of nitrogen and carbon release (Petersen, 1980; Sjursen and Holmstrup, 2004), the  
200 influence of the available resource quality (Chen et al., 1995) and the creation of  
201 nutrient sources for heterotrophic microbes and primary producers (Rusek, 1998).  
202 These processes should therefore be recognized as integral components of soil  
203 structure (Fjellberg, 1986; Rusek, 1985).

204 Indirect effects of microarthropods via the provision of organic material to  
205 microorganisms are not the only indirect mechanisms to be considered. The complex  
206 interactions with the larger components of the soil fauna have not yet been considered  
207 in detail in any study (but see Ponge 1988; Ponge, 1991; Salmon and Ponge, 2001). In  
208 our opinion, especially the interaction between different functional groups should be  
209 more closely investigated, as the biggest effect sizes are assumed to be found in this  
210 context rather than in studies dealing with direct effects.

211 Another important aspect is the impact of different agricultural practices on soil fauna  
212 and soil aggregation as abiotic factors. Once the biotic interactions between different  
213 faunal groups have become clearer, another focus should be on the impact of tillage,  
214 ploughing or compaction of soil on these interactions. It is known that different  
215 taxonomic groups respond differently to agricultural practices in different types of soils  
216 and depending on fertilizer additions etc. (see e.g. van Capelle et al., 2012, Roger-  
217 Estrade et al., 2010, for microarthropods see Ponge et al. 2013), however, closer  
218 investigation would be necessary in order to develop appropriate strategies to e.g.  
219 increase soil fertility and resistance towards erosional loss by increasing soil stability  
220 via soil fauna.

221 **Conclusion**

222 Despite their underrepresentation in the soil aggregation literature, we highlighted and  
223 discussed several potential mechanisms via which microarthropods could influence soil  
224 aggregation.

225 Due to their relatively small body size and total biomass, which is lower than that of  
226 fungi, bacteria and other taxa such as nematodes and protozoa, microarthropods may  
227 rather indirectly than directly affect soil structure. However, in some cases the impact  
228 of the production of assumedly large amounts of organic material in form of necromass,  
229 eggs, etc. might play an important role as direct starting points for microaggregate  
230 formation. We propose to start studying soil aggregation formation with easy-to-handle  
231 species such as *Folsomia candida* in experimental designs that allow assessing the  
232 direction and magnitude of the various possible mechanisms, especially direct vs  
233 indirect mechanisms. Difficulties with culturing microarthropods for experiments, but also  
234 with the collection of direct observations have hampered empirical studies to date. The  
235 usage of high resolution filming and photographing, which is nowadays very feasible  
236 given the remarkable advances in microscopy technologies, is necessary to observe  
237 how microarthropods act in the formation of soil aggregates. Coupling these  
238 technologies with small scale experimental designs will allow teasing apart the roles of  
239 various mechanisms that act simultaneously. An element of complexity and realism  
240 will be given by studies addressing the impact of different taxa (e.g. Collembola and  
241 Acari) on soil structure in opposition to studies focusing on species-specific effects. In  
242 this context, a focus should in our opinion be on the interaction of functionally  
243 defined, trait-based groups across all soil biota.

244

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251

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450 Figure captions

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452 Fig. 1: Overview of potential mechanisms used by microarthropods for the formation  
453 of soil aggregates. Mechanisms are divided into direct and indirect processes and  
454 based on Collembola and oribatid mites as most abundant soil microarthropod  
455 representatives.