

# Effect of Collembola on mineralization of litter and soil organic matter

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**Abstract** Although soil Collembola are known to contribute to soil carbon (C) cycling, their contribution to the mineralization of C sources that differ in bioavailability, such as soil organic C (SOC) and leaf litter, is unknown. Stable C isotopes are often used to quantify the effects of both soil C and litter C on C mineralization. Here, <sup>13</sup>C-labeled litter was used to investigate the effects of Collembola (*Folsomia candida*) on the mineralization of both SOC and litter C in laboratory microcosms. The three microcosm treatments were soil alone (S); soil treated with δ<sup>13</sup>C-labeled litter (SL); and soil treated with δ<sup>13</sup>C-labeled litter and Collembola (SLC). The presence of Collembola did not significantly affect soil microbial biomass or litter mass loss and only had a small effect on CO<sub>2</sub> release during the first week of the experiment, when most of the CO<sub>2</sub> was derived from litter rather than from SOC. Later, during the experiment (days 21 and 63), when litter-derived labile C had been depleted and when numbers of Collembola had greatly increased, Collembola substantially increased the emission of SOC-derived CO<sub>2</sub>. These results suggest that the effect of

Collembola on soil organic C mineralization is negatively related to C availability.

**Keywords** Collembola · Stable isotope · C mineralization · C availability

## Introduction

Collembola and other soil fauna play essential roles in the transformation of soil organic matter and in nutrient cycling (Rusek 1998). Although their direct contribution to energy flow is small, Collembola have a large indirect effect on the soil ecosystem (Petersen 2002; Ponge 2015) because they can regulate soil respiration, the leaching of dissolved organic carbon (C), and the mineralization of nitrogen (N) (Filser 2002; Ke et al. 2005; Lussenhop and BassiriRad 2005; Neher et al. 2012). However, the effects of Collembola on mineralization and stabilization of soil organic C (SOC) are poorly known. Microbial respiration and microbial biomass, for example, were increased (Addison et al. 2003; Coleman et al. 2004; Cragg and Bardgett 2001) or not affected by Collembola (Kaneda and Kaneko 2008; Kaneko et al. 1998).

According to the availability of C, the C pool in soil can be divided into stabilized C, potentially mineralizable C, and readily mineralizable C, and soil organisms can enhance the transformation of potentially mineralizable C into readily mineralizable C (Zhang et al. 2013). The contributions of soil fauna to C dynamics may be greatly affected by resource availability and/or quality. For instance, Yang and Chen (2009) reported that soil fauna enhanced soil organic C (SOC) decomposition to a greater extent in systems with less available resource (Yang and Chen 2009). Earthworms had a greater effect on the transformation of less labile C, such as SOC and litter C, than on the transformation of readily

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mineralizable C, such as the C in root exudates (Huang et al. 2015). Therefore, we wondered whether C availability also regulates the effects of Collembola on C transformations in soil. Given that labile C content may decrease and recalcitrant C content may increase over time as a C source incubates in soil, we hypothesized that Collembola would promote microbial activity and stimulate C mineralization of both litter C and SOC with the depleting of available C during an incubation experiment. We used a microcosm experiment to test this hypothesis in the current study. The use of labeled litter is a general method (Vidal et al. 2016, 2017) to distinguish between litter- and soil-derived C in emitted CO<sub>2</sub>. We chose the typical soil and typical leaf at an evergreen broad-leaf forest in subtropical China to evaluate the effect of Collembola on litter C and SOC mineralization.

## Materials and methods

### Soil, litter, and Collembola

Mineral soil, collected (0–10 cm) from a pine and broad-leaf mixed forest at the Dinghushan Biosphere Reserve (Guangdong Province, subtropical S China: 112° 10' E, 23° 10' N), is classified in the Ultisol group and Udult subgroup according to the USDA soil classification system (Huang et al. 2013); soil samples were air-dried, sieved (<2 mm) and sterilized in autoclaving at 121 °C for 30 min. The soil is a typical evergreen broad-leaf forest soil of subtropical China and had the following characteristics: pH 3.54, SOC 5.96%, and soil readily oxidized organic C (ROC after oxidation with 333 mM KMnO<sub>4</sub>) 1.31%.

Leaf litter of *Schima superba* (Gardner & Champ), a common plant of subtropical China, was used. *S. superba* seedlings were grown in pots in a growth chamber. When the seedlings were 50-cm high, they were transferred to greenhouse. From 12:00 hours to 13:00 hours every day for 7 days, <sup>13</sup>CO<sub>2</sub> (99 atom% <sup>13</sup>C, Cambridge Isotope Laboratories, Inc., USA) was pumped into the greenhouse atmosphere. Fully expanded leaves were then collected, oven-dried, and chopped into small pieces (2–3 mm), which were used as the litter in the experiment. The δ<sup>13</sup>C value of this litter was 1151.54‰. The method used to determine the δ<sup>13</sup>C value is described below. The Collembola *Folsomia candida* was cultured in the laboratory on dried yeast (Løkke 1998).

### Microcosms

Microcosms consisted of 150-ml beakers containing 100 g of soil (dry weight), which was dry when placed in the beakers. The 36 microcosms were placed in a plastic container (10.5-cm internal diameter × 12-cm height, one microcosm per container), that could be sealed with airtight lids. A 50-ml beaker

with 20 ml 1 M NaOH was also placed in each container which was sealed. The NaOH was used to adsorb evolved CO<sub>2</sub> (see “CO<sub>2</sub> emission” section). To inoculate the soil with an assemblage of microorganisms, 30 g of fresh soil were mixed with 300 ml of sterile water. After most of the soil particles had settled, the supernatant was passed through a 23-μm sieve to permit the passage of microorganisms; then, 10 ml were added to each microcosm. An additional 30 ml of sterile water were then added to each microcosm to increase the water content to 60% of maximum holding capacity. The soil water content was maintained constant by weighing each beaker and adding sterile water as needed every other week. A 1-g quantity of labeled litter (δ<sup>13</sup>C = 1151.54‰) was placed on the soil surface of each of 24 of the 36 microcosms. Thirty mature Collembola (*F. candida*) were then added to 12 of the microcosms that contained litter. The experiment included three treatments: soil alone (S), soil treated with δ<sup>13</sup>C-labeled litter (SL), and soil treated with δ<sup>13</sup>C-labeled litter and Collembola (SLC). Each treatment was replicated 12 times, microcosms were incubated in a climate chamber under the darkness at 25 °C. Four replicate microcosms of each treatment were destructively sampled at 7, 21, and 63 days.

### Analyses

#### CO<sub>2</sub> emission

The CO<sub>2</sub> evolved from each microcosm was trapped in the 20 ml 1 M NaOH. Every 2 weeks, the NaOH was exchanged and the trapped CO<sub>2</sub> was measured by titration with 0.5 M HCl after carbonate was precipitated with 5 ml saturated BaCl<sub>2</sub> solution. To determine the δ<sup>13</sup>C values of carbonate, the solution with the precipitate was transferred to 50-ml centrifuge tubes. The tubes were shaken by hand and centrifuged for 2 min at 2500×g. After centrifugation, the supernatant was discarded, and the tubes with pellets of BaCO<sub>3</sub> were immediately filled again with H<sub>2</sub>O and then closed, shaken, and centrifuged (2500×g). This washing procedure was repeated until the pH of the supernatant was 6.8–7.0, thereby preventing further trapping of CO<sub>2</sub> from the air. The samples were then dried in an oven at 80 °C for 48 h (Marhan et al. 2007); δ<sup>13</sup>C values were measured with an IsoTope 100 stable isotope ratio gas mass spectrometer, IsoPrime, UK.

#### Sampling of soil, litter, and Collembola

Before a microcosm was destructively sampled, the litter was carefully removed from the soil surface, dried at 60 °C, and then pulverized in a ball mill. We considered a homogenous distribution of Collembola in the microcosms; the soil of each microcosm was then vertically divided into two parts: one part was used for extraction of Collembola, and the other part was used for analysis of microbial phospholipids fatty acids (PLFAs), SOC,

and ROC. Collembola were extracted using Tullgren dry funnels for 48 h; the extracted specimens were stored in alcohol and then counted. PLFAs were analyzed as described by Bossio and Scow (1998). Concentrations of each PLFA were standardized relative to the 19:0 internal reference concentration. Bacterial biomass was represented by i14:0, i15:0, i16:0, a15:0, i17:0, a17:0, 15:0 3OH, 16:1 2OH, 16:1ω5c, 16:1ω9c, 18:1ω5c, 18:1ω7c, 18:1ω9c, cy17:0, cy19:0, 14:0, 15:0, 17:0, and 18:0 and fungal biomass was represented by 18:2ω6,9 and 18:3ω6,9,12c (Baath 2003; Bossio and Scow 1998; Frostegard et al. 2011). The ratio of cy17:0 to 16:1ω7c was used to indicate the activity of bacteria, with the higher ratio values indicating bacterial stress (Grogan and Cronan 1997). SOC and ROC were analyzed by the  $K_2Cr_2O_7$  oxidation-external heating method and by the  $KMnO_4$  oxidation method, respectively (Blair et al. 1995, 1997).

*Identification of the C sources in released CO<sub>2</sub> using stable isotopes*

When *n* isotope systems are used to determine the proportional contributions of *n* + 1 sources to a mixture, standard linear mixing models can be used to mathematically solve for the unique combination of source proportions that conserves mass balance for all *n* isotopes (Phillips and Gregg 2003). In our study, with one isotope system and two sources (C from litter and C from soil organic carbon), the following mass balance equation can be solved to determine the proportions ( $f_{litter}, f_{soil}$ ) of source isotopic signatures ( $\delta_{litter}, \delta_{soil}$ ) that coincide with the observed signature as the

$$\delta_{CO_2} = f_{litter} \times \delta_{litter} + f_{soil} \times \delta_{soil} \tag{1}$$

and

$$f_{litter} + f_{soil} = 1 \tag{2}$$

where  $f_{litter}$  and  $f_{soil}$  refers to the proportion of litter-derived and soil-derived CO<sub>2</sub>, respectively; and  $\delta_{litter}$  and  $\delta_{soil}$  refers to the isotope abundance ( $\delta^{13}C$ ) of litter and soil, respectively.

**Statistics**

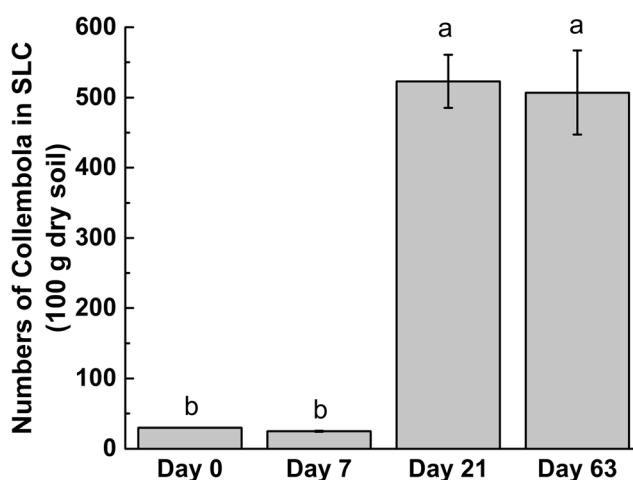
To assess the effect of Collembola on mineralization of litter, mineralization of SOC, and soil microbial PLFAs, we used a one-way ANOVA, with litter dry weight being the dependent variable, and repeated measures ANOVAs, with ROC, PLFAs, CO<sub>2</sub>, and  $\delta^{13}C$  values of CO<sub>2</sub> being the dependent variables. When ANOVA gave significant differences, we also determined differences among means using LSD post hoc multiple comparisons. The significance level in all analyses was  $P < 0.05$ . All data passed the homogeneity of variance test, and all statistical analyses were carried out using SPSS 16.0. Figures were drawn with Origin 2016.

**Results**

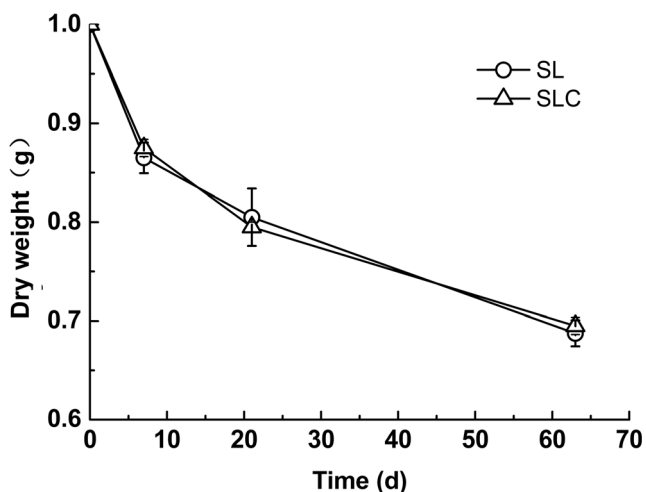
Collembola numbers were about 17-fold higher on days 21 and 63 than on days 0 and 7 (Fig. 1). By day 63, litter mass on the soil surface had declined by about 33% but the decline was unaffected by Collembola ( $F = 0.011, P = 0.918, \text{Fig. 2}$ ). During the whole experiment, the C content of the litter did not change significantly ( $F = 0.832, P = 0.489, \text{Fig. 3a}$ ), whereas the N content of the litter significantly increased from  $1.76 \pm 0.07\%$  on day 0 to  $2.66 \pm 0.41\%$  on day 63 ( $F = 105, P < 0.01, \text{Fig. 3b}$ ). There were no significant differences in C or N contents of litter (% C:  $F = 0.005, P = 0.947$ ; % N:  $F = 1.138, P = 0.297, \text{Fig. 3}$ ), or in C/N ratio of litter between microcosms with or without Collembola ( $F = 1.126, P = 0.299, \text{Fig. 4}$ ).

Bacterial biomass did not significantly differ among the three treatments during the incubation ( $F = 1.944, P = 0.205, \text{Fig. 5a}$ ). Although fungal biomass differ significantly at different sampling time ( $F = 7.177, P = 0.014, \text{Fig. 5b}$ ) it did not differ significantly among the three treatments (day 7:  $F = 1.291, P = 0.321$ ; day 21:  $F = 0.641, P = 0.549$ ; day 63:  $F = 1.138, P = 0.362$ ; Fig. 5b). At any sampling time, soil microbial biomass did not significantly differ among the three treatments ( $F = 0.312, P = 0.740$ ; Fig. 5c), but ratio of fungal biomass to bacterial biomass differ significantly at different sampling time ( $F = 6.024, P = 0.030, \text{Fig. 5d}$ ). The ratio of cy17:0 to 16:1ω7c did not differ among treatments during the incubation ( $F = 0.897, P = 0.441$ ; Fig. 6).

The presence of Collembola and litter did not significantly affect the SOC content of microcosms (data not shown) ( $F = 0.591, P = 0.547$ ). The ROC content did not differ among treatments on day 7 but it did differ among treatments on day



**Fig. 1** Collembola numbers in soil microcosms, containing  $\delta^{13}C$ -labeled litter (treatment SLC) as affected by incubation time. This is an estimation value according to the number of Collembola in a part of soil of the microcosm. Values are means  $\pm$  SE ( $n = 4$ ). Means with the different letters are significantly different ( $P < 0.05$ )



**Fig. 2** Changes in litter mass remaining on the soil surface of soil treated with  $\delta^{13}\text{C}$ -labeled litter (SL); and soil treated with  $\delta^{13}\text{C}$ -labeled litter and Collembola (SLC). Values are means  $\pm$  SE ( $n = 4$ )

21 ( $F = 158$ ,  $P < 0.01$ ), when the ROC content was significantly increased in S microcosms, unchanged in SL microcosms and decreased in SLC microcosms (Fig. 7). ROC content did not differ among treatments on day 63, but the ROC content increased when litter was added and again when litter and Collembola were added. ROC content in SL and SLC microcosms were significantly higher on day 63 than on day 21.

The cumulative release of  $\text{CO}_2$  showed significant difference among treatments ( $F = 96$ ,  $P < 0.01$ , Table 1). Early during the incubation,  $\text{CO}_2$  emission rate was much greater in SL and SLC microcosms than in S microcosms (Fig. 8).  $\text{CO}_2$  emission rate did not significantly differ between SL and SLC microcosms before day 35 but then it was higher in SLC microcosms than in SL microcosms, perhaps because numbers of Collembola had greatly increased during this period.

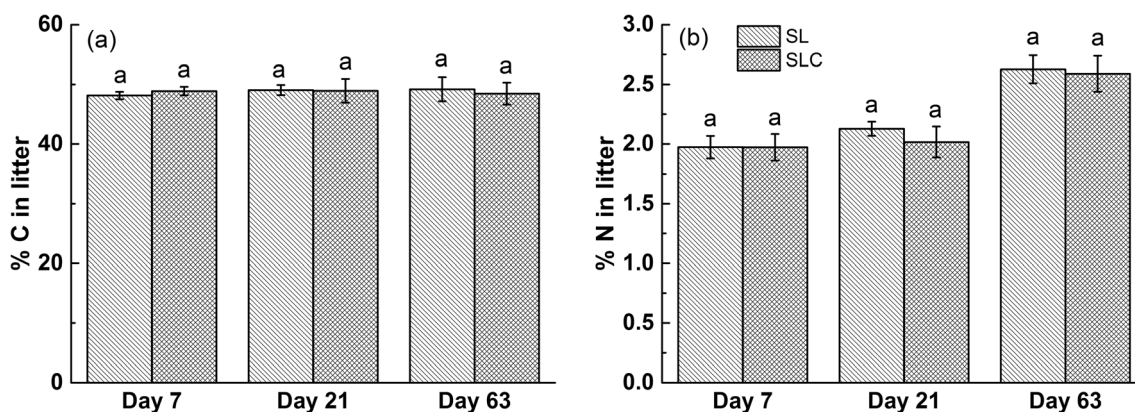
The  $\delta^{13}\text{C}$  values of  $\text{CO}_2$  were significantly higher in SL and SLC microcosms than S microcosms but this difference gradually decreased over time ( $F = 1012.14$ ,  $P < 0.01$ , Fig. 9).

The  $\delta^{13}\text{C}$  values of  $\text{CO}_2$  did not significantly differ between SL and SLC microcosms at days 7 to 21. But the  $\delta^{13}\text{C}$  values was lower in SLC than in SL microcosms from day 35 to the end of experiment and this suggests that increases in numbers of Collembola were associated with a decrease in the proportion of litter-derived C in the  $\text{CO}_2$  emission.

According to the amount (mg of C) of  $\text{CO}_2$  and  $f_{\text{litter}}$  and  $f_{\text{soil}}$  values, we calculated the amounts (mg of C  $\text{day}^{-1}$ ) of litter- and soil-derived  $\text{CO}_2$  and the net effect of Collembola (Table 2). This indicated that soil organisms other than Collembola mineralized the litter-derived C and that most of the litter-derived  $\text{CO}_2$  was generated by soil organisms other than Collembola. In other words, Collembola had little effect on litter-derived  $\text{CO}_2$ . The net contribution of Collembola to soil-derived  $\text{CO}_2$  was substantial on day 63 (Table 2). In others, the net effect of Collembola to the mineralization of soil C was about 0 on day 7 but increased to 26% on day 21 and to 24.16% on day 63. In other words, Collembola substantially increased the emission of soil-derived C with the incubation time.

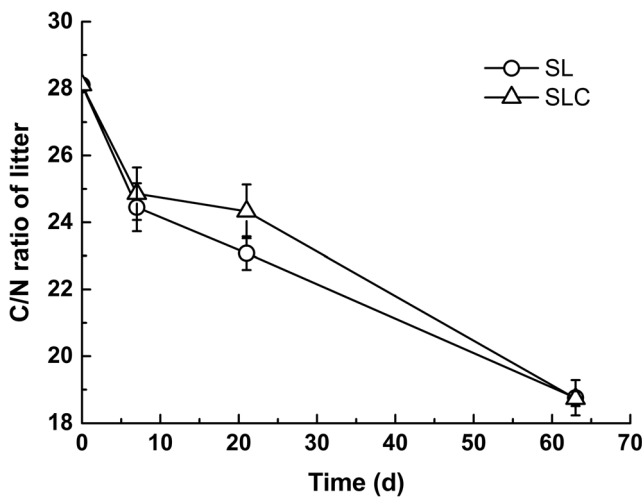
## Discussion

In our experiment, the Collembola *F. candida* had no significant effect on the C and N contents and thus to C/N ratio in the microcosms. The Collembola also had no effect on litter mass loss or litter C mineralization. These results are consistent with previous studies (Andren and Schnurer 1985; Aslam et al. 2015; Chamberlain et al. 2006). Because *F. candida* is a fungivorous (Chahartaghi et al. 2005; Haubert et al. 2006; Scheu and Simmerling 2004), it does not directly assimilate C from litter but indirectly affects litter decomposition by feeding on microorganisms and by altering the soil microenvironment (Addison et al. 2003; Filser 2002). In other studies, however, mechanical fragmentation of litter, which was done in the current research, facilitated the interaction between



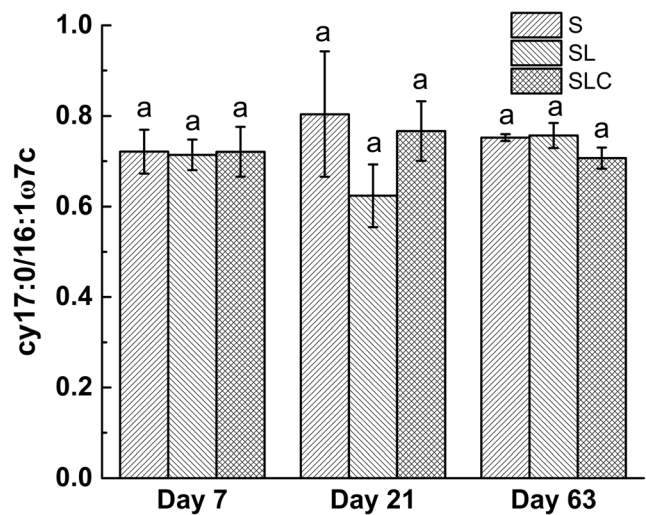
**Fig. 3** The C (a) and N contents (b) of litter of soil treated with  $\delta^{13}\text{C}$ -labeled litter (SL); and soil treated with  $\delta^{13}\text{C}$ -labeled litter and Collembola (SLC). Values are means  $\pm$  SE ( $n = 4$ ). Means with the different letters are significantly different ( $P < 0.05$ )





**Fig. 4** Changes in C/N ratio of litter of soil treated with  $\delta^{13}\text{C}$ -labeled litter (SL) and soil treated with  $\delta^{13}\text{C}$ -labeled litter and Collembola (SLC). Values are means  $\pm$  SE ( $n = 4$ )

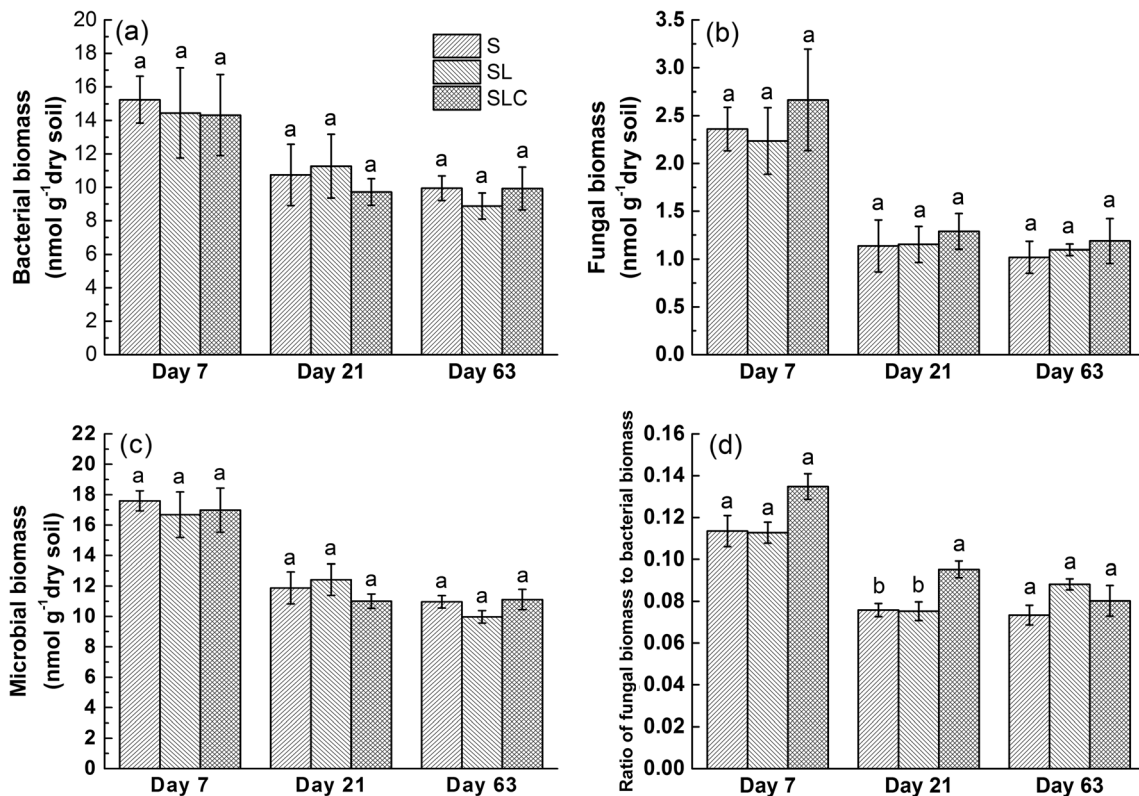
Collembola and microorganisms, which in turn accelerated litter decomposition (Hunter et al. 2003; Pieper and Weigmann 2008; Seeber et al. 2006; Yang et al. 2012). However, the litter used in our study was not obtained from senescence leaves. The nutrient content of the litter was higher than the dead leaves, especially the content of phosphorus (Gusewell and Verhoeven 2006; Liu et al. 2007). The



**Fig. 6** The ratio of cy17:0 to 16:1 $\omega$ 7c in soil (S); soil treated with  $\delta^{13}\text{C}$ -labeled litter (SL); and soil treated with  $\delta^{13}\text{C}$ -labeled litter, and Collembola (SLC). Values are means  $\pm$  SE ( $n = 4$ ). Means with the different letters are significantly different ( $P < 0.05$ )

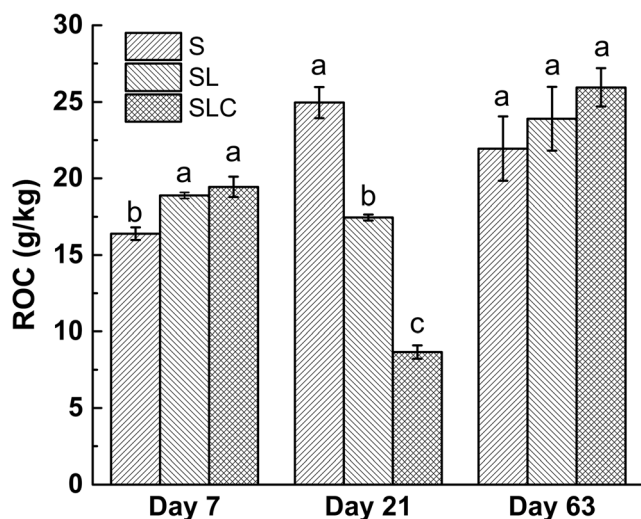
difference on litter may explain these differences. Thus, whether Collembola indirectly accelerate litter decomposition this might depend on litter quality and Collembola species.

We also found that  $\text{CO}_2$  emissions at day 7 were mainly derived from the litter. In previous reports, C was mineralized more rapidly from litter than from SOC (Kuzyakov 2011;



**Fig. 5** Bacterial biomass (a), fungal biomass (b), total microbial biomass (c), and ratio of fungal biomass to bacterial biomass (d) in soil (S); soil treated with  $\delta^{13}\text{C}$ -labeled litter (SL); and soil treated with  $\delta^{13}\text{C}$ -labeled

litter, and Collembola (SLC). Values are means  $\pm$  SE ( $n = 4$ ). Means with the different letters are significantly different ( $P < 0.05$ )



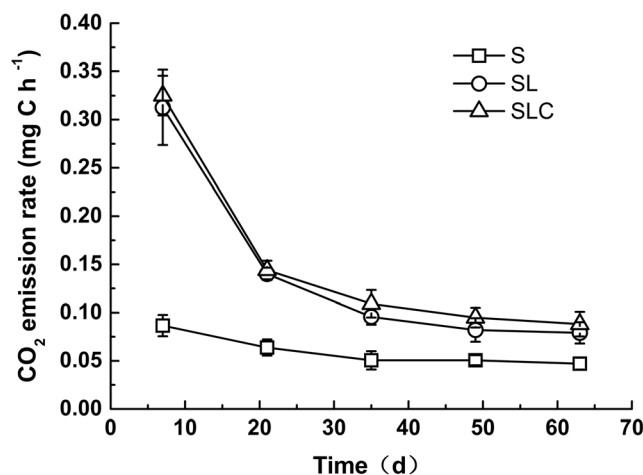
**Fig. 7** The content of readily oxidized organic C (ROC) in soil (S); soil treated with  $\delta^{13}\text{C}$ -labeled litter (SL); and soil treated with  $\delta^{13}\text{C}$ -labeled litter, and Collembola (SLC). Values are means  $\pm$  SE ( $n = 4$ ). Means with the different letters are significantly different ( $P < 0.05$ )

Pausch and Kuzyakov 2012; Schlesinger and Lichter 2001). Because our results indicated that litter decomposition was not directly affected by *F. candida*, we infer that most of the litter mass loss and C mineralization was caused by microbial decomposition of the available C in the litter.

Along with mites, Collembola constitute an important mesofauna component of almost all terrestrial ecosystems (Rusek 1998). Collembola can increase nutrient cycling, improve soil physical and chemical characteristics, regulate soil microbial composition, and enhance soil fertility (Chen et al. 2007). However, their direct contribution to energy flow is limited and they mostly appear to have indirect effects on soil processes (Petersen 2002). In previous studies, for example, Collembola affected soil C turnover mainly by affecting soil structure and microbial activity (Addison et al. 2003; Cragg and Bardgett 2001; Endlweber et al. 2009; Lavelle 1997). In our study, Collembola had no significant effect on the total C pool in soil, but they affected ROC and the relative quantities of C in the different C pools, and also enhanced the mineralization of soil-derived C. In SL microcosms on day 21, soil microorganisms may have preferentially used litter C rather than SOC. In SLC microcosms on day 21, litter C was also preferred by soil microorganisms and Collembola;

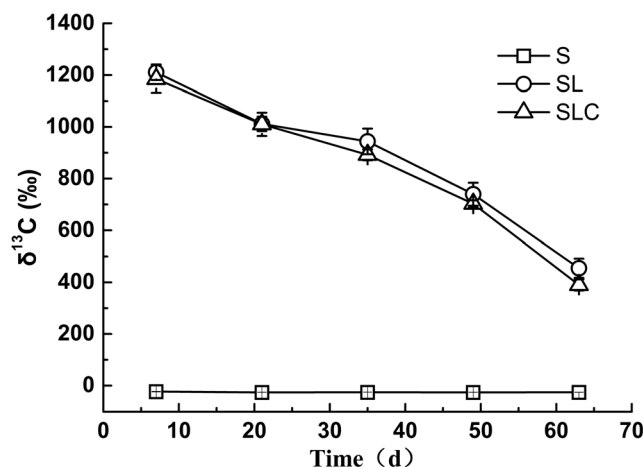
**Table 1** Cumulative  $\text{CO}_2$  respired from microcosms over the 63-day experimental period. Values are means (SE) ( $n = 4$ ). Different letters indicate significant differences ( $P < 0.05$ )

Microcosm	Cumulative $\text{CO}_2$ (mg C- $\text{CO}_2$ )
S (soil)	86.0 (10.5)b
SL (soil and litter)	181.0 (13.4)a
SLC (soil, litter, and Collembola)	199.1 (12.9)a



**Fig. 8** Changes in  $\text{CO}_2$  emission rate in soil (S); soil treated with  $\delta^{13}\text{C}$ -labeled litter (SL); and soil treated with  $\delta^{13}\text{C}$ -labeled litter, and Collembola (SLC). Values are means  $\pm$  SE ( $n = 4$ )

meanwhile, the numbers of Collembola had greatly increased and may have mineralized soil-derived C and thereby decreased ROC. At day 63, when the labile C in soil had been depleted and Collembola numbers were also high, Collembola would transfer soil C from stable to more labile pools. Thus, the increased content of ROC occurred at day 63 when Collembola numbers were high, suggesting that the Collembola may have increased the release of soil-derived C by increasing the pool of labile C in soil. Because Collembola promoted the release of soil-derived C, they reduced the  $\delta^{13}\text{C}$  values of  $\text{CO}_2$  from days 35 to 63 (Fig. 9). Previous studies (Fox et al. 2006; Johnson et al. 2005) also found that Collembola enhanced the mineralization of soil C. For example, Fox et al. (2006) confirmed that cumulative  $\text{CO}_2$  emissions from soil were significantly increased when Collembola were present and that this resulted from the transfer of soil C from stable to more labile pools.



**Fig. 9** The  $\delta^{13}\text{C}$  values of  $\text{CO}_2$  in soil (S); soil treated with  $\delta^{13}\text{C}$ -labeled litter (SL); and soil treated with  $\delta^{13}\text{C}$ -labeled litter, and Collembola (SLC). Values are means  $\pm$  SE ( $n = 4$ )

**Table 2** Effects of Collembola on litter- and soil-derived CO<sub>2</sub> (mg C day<sup>-1</sup>) on days 7, 21, and 63. SL microcosms contain soil and δ<sup>13</sup>C-labeled litter. SLC microcosms contain soil, δ<sup>13</sup>C-labeled litter, and Collembola. Values are means (SE) (n = 4)

	CO <sub>2</sub> litter			CO <sub>2</sub> soil		
	Day 7	Day 21	Day 63	Day 7	Day 21	Day 63
Microcosm						
SL (with litter)	7.85 (0.36)	2.84 (0.31)	0.79 (0.38)	0 (0)	0.37 (0.18)	1.13 (1.00)
SLC (with litter and Collembola)	8.04 (0.36)	2.84 (0.74)	0.80 (0.57)	0 (0)	0.50 (0.26)	1.49 (0.47)
Net effect of Collembola (%)	2.36 (0.05)	0 (0)	1.25 (0.85)	0 (0)	26 (1.5)	24.16 (0.24)

In this study, we did not measure the respiration of Collembola, and thus could not quantify the contribution of Collembola to C mineralization. Some studies had shown that Collembola respiration accounts for 1% of total soil respiration (Coulson and Whittaker 1978), while Collembola respiration may not be ignored when the population size of Collembola, e.g., in this study, was much greater than those in common soil situations. Thus, we made a rough calculation of the Collembola respiration according to dataset from literatures (Lagerlöf and Andrén 1985; Persson et al. 1980). However, the estimated respiration of Collembola was even bigger than the total soil respiration. So, we are wondering whether this method of respiration estimation of Collembola is suitable to our laboratory experiment. Another laboratory experiment found that CO<sub>2</sub> flux did not differ between soils with and without abundant Collembola (5 individuals g<sup>-1</sup> dry soil) (Chamberlain et al. 2006). Therefore, it was also possible that the respiration of a large-sized population of Collembola still accounted for a small part of the total respiration. Nevertheless, the aim of this study is to evaluate the effects of Collembola presence on C mineralization, rather than an estimation of Collembola respiration. It suggests that total C mineralization was controlled by C availability and conversion of soil C was enhanced by the soil food web. Litter addition at the start of our experiment increased microbial respiration rapidly probably due to the rapid mineralization of the labile C of the litter (Fig. 8). Also, during the early stages of incubation, CO<sub>2</sub> emissions were derived mostly from litter C rather than from SOC (Table 2). Later, during the incubation, however, C mineralization depended more on SOC than on litter C, especially when Collembola were present. Collembola can affect soil C processes by affecting soil physical, chemical, and microbiological properties (Addison et al. 2003; Filser 2002). Indeed the effects of Collembola on C and N mineralization may be mediated by their effects on microbial community composition (Kaneda and Kaneko 2008; Seastedt 1984; Teuben 1991).

## Conclusions

Collembola significantly increased the mineralization of SOC at the late stage of incubation when litter labile C was

depleted. There was no evidence that Collembola affected the litter decomposition rate or microbial biomass. In this study, we only focused on the effects of Collembola (*F. candida*) on the mineralization of SOC and leaves of an evergreen broad-leaf forest in subtropical China. Future studies should investigate the effects of Collembola on C processes in different types of soil and litter.

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## Compliance with ethical standards

**Conflict of interest** The authors declare that they have no conflict of interest.

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