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Simulated leaf litter addition causes opposite priming effects on natural forest and plantation soils

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Abstract

The conversion of natural forests to tree plantations alters the quality and decreases the quantity of litter inputs into the soil, but how the alteration of litter inputs affect soil organic matter (SOM) decomposition remain unclear.We examined SOM decomposition by adding ¹³C-labeled leaf-litter of Chinese fir (*Cunninghamia lanceolata* (Lamb.) Hook) to soils from a natural evergreen broadleaved forest and an adjacent Chinese fir plantation converted from a natural evergreen broad-leaved forest 42 years ago. Over 195 days, we monitored CO₂ efflux and its $\delta^{13}C$, microbial biomass, and the composition of microbial groups by phospholipid fatty acids (PLFAs). To distinguish priming mechanisms, partitioning of C sources in CO₂ and microbial biomass was determined based on δ^{13} C. Leaf-litter addition to natural forest increased microbial biomass and induced up to 14% faster SOM decomposition (positive priming) than that in soil without litter. In contrast, negative priming in soils under plantation indicated preferential use of added leaf-litter rather than recalcitrant SOM. This preferential use of leaf-litter was supported by an increased fungal to bacterial ratio and litter-derived (^{13}C) microbial biomass, reflecting increased substrate recalcitrance, the respective changes in microbial substrate utilization and increased C use efficiency. The magnitude and direction of priming effects depend on microbial preferential utilization of new litter or SOM. Concluding, the impact of coniferous leaf-litter inputs on the SOM priming is divergent in natural evergreen broad-leaved forests and plantations, an important consideration in understanding long-term C dynamics and cycling in natural and plantation forest ecosystems.

Keywords Microbial community composition . Priming effects . Selective decomposition . Subtropical forest soils . Land-use effects

Introduction

Identifying land-use and management practices that maintain and increase soil carbon (C) storage is important for C

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removal from the atmosphere and improving ecosystem services (Herrero et al. [2016\)](#page-8-0). There is debate about the amount and duration of additional C that can be stored annually in soil following changes in land use or management (Smith [2014\)](#page-9-0).

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Land-use changes from natural to artificial ecosystems, e.g., plantations, in tropical and subtropical areas contribute 12– 15% of the global anthropogenic $CO₂$ emissions (Van der Werf et al. [2009](#page-9-0); Pan et al. [2011\)](#page-9-0). The tropics and subtropics have the largest area of plantation forests worldwide (Brockerhoff et al. [2008;](#page-8-0) Huang et al. [2013](#page-8-0)).

Approximately 65% of the Chinese forests and associated C stocks are located in the southeastern subtropical provinces of Fujian, Hunan, and Zhejiang (Piao et al. [2009](#page-9-0)). More than 32% of these forests are plantations with the dominant species being Chinese fir (Cunninghamia lanceolata) (see Fig. S1; FAO [2006\)](#page-8-0). Most of these plantations were converted from natural evergreen broad-leaved forests over the past decades (SFA [2009](#page-9-0); Lu et al. [2015\)](#page-8-0). Such ecosystem conversion from diverse broad-leaved forests to conifer monocultures has resulted in decreased litter quality and quantity that influences soil C storage, decomposition, and dynamics (Chen et al. [2005;](#page-8-0) Yang et al. [2009](#page-9-0)). The litter from Chinese fir has higher ratios of C to nitrogen (N) and lignin to N than those of the broad-leaved tree species (Lin et al. [2011](#page-8-0)), and therefore, a slower decomposition rate (Yu et al. [2015\)](#page-9-0).

In this context, the priming effect (PE) is defined as the acceleration or retardation of SOM decomposition, i.e., strong short-term changes in soil organic matter (SOM) decomposition caused by fresh C inputs (Kuzyakov et al. [2000;](#page-8-0) Kuzyakov [2010\)](#page-8-0), which can accelerate SOM decomposition (i.e., positive PE) up to 380% or decrease SOM decomposition (i.e., negative PE) up to 50% (Kuzyakov et al. [2000;](#page-8-0) Kuzyakov [2002](#page-8-0); Cheng et al. [2014](#page-8-0)). Current views suggest that PE mainly depends on changes in microbial activities and/or strategies after fresh substrate inputs, which provide easy-to-use nutrient and/or more energy for microbial growth (Fontaine et al. [2003](#page-8-0); Blagodatskaya and Kuzyakov [2008\)](#page-8-0). However, the extent to which priming changes SOM dynamics could be leveled out by microbial adaptation to substrate decomposition. This suggests that microorganisms adapt to the litter, which decomposes faster when placed in its habitat of origin than in other environments (Ayres et al. [2009](#page-8-0); Veen et al. [2015;](#page-9-0) Yu et al. [2015](#page-9-0)). Consequently, any changes in the response of soil-C decomposition to the litter addition could be ascribed to the adaptation of the microbial community (Cheng [1999;](#page-8-0) Meidute et al. [2008](#page-8-0); Lü et al. [2015;](#page-8-0) Don et al. [2017\)](#page-8-0). Negative PE occurs when microorganisms utilize the newly added substrate more than the existing SOM. Positive PE may occur when the added substrate induces microbial growth and activity including extracellular enzyme production (Blagodatskaya et al. [2014\)](#page-8-0), and therefore, microorganisms utilize SOMmoreintensively (Kuzyakov et al. [2000\)](#page-8-0). The above mechanisms might be important determinants of the SOM fate when tree species dominance changes, and especially after conversion from a natural forest to monoculture plantation.

Here, we present a case study based on a lab incubation of soils from two adjacent stands (one natural forest and one

converted to a Chinese fir plantation), to understand the consequences of the input of Chinese fir litter to SOM decomposition. By using the same litter, any changes in the response of soil-C decomposition to the litter addition are directly or indirectly linked to the nutrient status (e.g., N) and microbial community. We hypothesized that (1) the addition of leaf-litter from the Chinese fir plantation would stimulate microbialmediated SOM decomposition in natural forest soils (i.e., positive PE) because these microorganisms are not adapted for this specific litter, whereas (2) similar inputs of leaf-litter to plantation soil would facilitate the decomposition of litterderived C instead of SOM mineralization (i.e., negative PE) owing to the microbial adaptation of litter decomposition and consequently, faster and more complete litter processing. We base this second hypothesis on the fact that the coniferous plantation soils have higher fungal abundance compared with natural forest soils (Guo et al. [2016\)](#page-8-0).

Materials and methods

Soil sampling

In March 2015, soils were sampled from two forests: a mature natural forest dominated by Castanopsis kawakamii Hayata (Fagaceae) and an adjacent 42-year-old firstrotation Chinese fir plantation (converted from the above natural forest in 1973; hereinafter referred to as plantation), located at the Forest Ecosystem and Global Change Research Station (FEGCRS) (26°09′24″ N, 117°28'03″ E, 300 a.s.l.), Sanming (Lyu et al. [2017](#page-8-0); Liu et al. [2017\)](#page-8-0). Four replicate plots $(20 \times 20 \text{ m})$ were randomly selected in each forest type. Each plot was separated by buffer zones of more than 20 m. In each plot, 20 points were randomly selected to collect soil samples (top 10 cm) with an auger (3.5-cm diameter). The 20 soil cores were then pooled as a single representative sample from each plot. In total, four composite soil samples were collected from each forest type. The soil is classified as latosol soil in the Chinese soil classification, equivalent to Oxisol in the USDA Soil Taxonomy (State Soil Survey Service of China [1998](#page-9-0); Guo et al. [2016](#page-8-0)). General soil characteristics are described in Table [1.](#page-2-0) As this study is unreplicated at the stand level, it is best interpreted as a case study of stand differences which result from the conversion to plantation 42 years ago, though no treatment data or replicate stands are available.

Pulse labeling of Chinese fir seedlings

Pulse labeling was performed in three-quarters of the Chinese fir seedlings from July 18, 2014, to January 26, 2015, when the seedlings were about 1 m high. The remaining unlabeled seedlings were used to measure natural ¹³C abundance. Each

Table 1 Soil properties under natural evergreen broad-leaved forest and Chinese fir plantation

Property	Natural forest	Plantation	
pH	4.31 ± 0.02	3.72 ± 0.02	
Sand $(\%)$	34.7 ± 1.5	36.5 ± 2.8	
Clay $(\%)$	38.8 ± 0.1	37.7 ± 0.6	
Silt $(\%)$	26.5 ± 0.8	25.8 ± 0.4	
SOC content $(g \text{ kg}^{-1})$	29.2 ± 0.4	26.7 ± 0.5	
$\delta^{13}C$ (%o)	-27.9 ± 0.1	-26.5 ± 0.1	
N content (g kg^{-1})	2.03 ± 0.05	1.91 ± 0.03	
CN ratio	14.4 ± 0.3	14.0 ± 0.1	
NH_4^+ (mg N kg ⁻¹) ^a	47.3 ± 1.1	29.0 ± 0.5	
$NO_3^- (mg N kg^{-1})^a$	12.5 ± 0.6	14.6 ± 0.4	

Results are means \pm standard errors (*n* = 4)

 ${}^{\text{a}}$ NH₄⁺ and NO₃⁻ was measured once before incubation in March 2015

seeding was moved into a chamber $(2 \times 1.8 \times 1.5 \text{ m})$ for 13^1 CO₂ pulse labeling twice monthly (Fig. S2). The 13^1 CO₂ pulse was started from 8:00 to 12:00 a.m. by slowly injecting 20 ml 0.1 M HCl with a syringe into a bottle containing dissolved 13 C-sodium carbonate (99% 13 C, Cambridge Isotope Laboratories Inc., Andover, MA, USA). A total quantity of 5 g $NaH¹³CO₃$ was used every time for each chamber. The released ${}^{13}CO_2$ was pumped into the chamber, which had a fan to homogenize the internal air. A pulse was given once per hour for 4 h. Seedlings were immediately removed from the chamber after being labeled. Following labeling (denoted as day 0), the Chinese fir leaf-litter was cut with scissors in both unlabeled and labeled plots.

Laboratory incubation

After removing roots and stones, each soil sample was sieved (< 2 mm) while still field-moist. The sieved soil samples were stored at 4 °C until the incubation experiment started (about 2 weeks). Each replicate of each composite soil sample was incubated with either the addition of ¹³C-labeled (≈ + 150‰) Chinese fir leaf-litter (litter addition) or no addition (control). The litter C and N content and C/N ratio were 45.2%, 0.83%, and 54.5, respectively.

Forty grams (oven-dry equivalent) of soil were weighed into 50-ml polyvinylchloride cups, which were placed into a 1000-ml Mason jar and pre-incubated for 7 days at 20 °C with a constant soil moisture at 50% of the field waterholding capacity (determined on a separate sample). After the pre-incubation, 13C-labeled litter was mixed with soil homogeneously at a rate of 675 mg C of 13 C-labeled litter per kilogram soil (calculated based on the annual litter fall of Chinese fir plantations and the bulk density of the top 10 cm of soils). Soil moisture was adjusted with deionized water and constantly maintained at 60% of the waterholding capacity for the whole incubation period. This soil moisture was close to average field conditions and ensured that microbial processes were not water limited.

Measurement of $CO₂$ fluxes and isotope ratios

We used the alkali absorption method to trap evolved $CO₂$ from soils. A glass vial containing 20 mL NaOH solution (0.5 M) was placed in each Mason jar to trap $CO₂$ released from the incubating soil. Jars were kept sealed. Three Mason jars without soil were used to trap $CO₂$ from the ambient air trapped in each jar. The $CO₂$ captured by the NaOH solution was measured after 1, 4, 8, 14, 22, 31, 51, 74, 103, 133, and 195 days by titration with 0.25 M HCl after adding 5 mL 0.5 M BaCl₂ to precipitate all carbonates. The solution was filtered to separate the $BaCO₃$ precipitate, which was then heated at 50 \degree C for 24 h. Soil CO₂ production was calculated from the difference between the values of evolved $CO₂$ in the Mason jars minus the mean of those without soil. The δ^{13} C was measured using a GasBench II coupled to an isotope ratio mass spectrometer (Finnigan MAT-253, Thermo Electron). The δ^{13} C of soil samples was measured with a continuous flow isotope ratio mass spectrometer. The standard deviation of δ^{13} C often repeated samples was $< 0.3\%$ o.

Analyses of soil properties, microbial biomass, and main microbial groups

Soil C and N concentrations were determined using a Vario MAX CN elemental analyzer (Elementar Vario EL III, Germany). Soil texture was measured using a Mastersizer 2000 particle-sizing instrument (Malvern Instruments, Worcestershire, UK). Soil pH was measured with a pH meter, in a 1:2.5 mass/volume soil and water suspension. Soil availability N was extracted with 2 M KCl to determine the exchangeable NH_4^+ -N and NO_3^- -N concentrations, using a SKALAR San⁺⁺ Analyzer (Skalar, Breda, The Netherlands).

Destructive samplings of soils from four replicates per treatment were performed at 0, 30, 133, and 195 days to determine the concentrations of dissolved organic C (DOC) and microbial biomass C (MBC), which was measured by the chloroform fumigation extraction method (Vance et al. [1987\)](#page-9-0). At the end of incubation, the composition of microbial functional groups was evaluated using phospholipid fatty acid (PLFA) analysis (Tunlid et al. [1989](#page-9-0); Lü et al. [2015;](#page-8-0) Gunina et al. [2014](#page-8-0)).

Partitioning of C sources in $CO₂$ and microbial biomass and calculation of priming effects

The ¹³C labeling of Chinese fir leaf-litter allowed the separation of soil (Rs) and leaf-litter-derived C (Rc) in released $CO₂$ using the mass balance approach. Priming effect (μ g C g⁻¹) of leaf-litter addition on native SOM mineralization was defined

by the following equations, using the symbols proposed by Fontaine et al. [\(2011](#page-8-0)):

$$
Rs + RC = Rt \tag{1}
$$

$$
Rs \times As^{13} + Rc \times Ac^{13} = Rt \times At^{13}
$$
 (2)

where As^{13} is the ¹³C abundance of soil C, Ac^{13} is the ¹³C abundance of leaf litter. *Pt* is the total CO, emitted from soil abundance of leaf-litter, Rt is the total $CO₂$ emitted from soil with leaf-litter, and At^{13} is ¹³C abundance measured in the CO₁ tran. $CO₂$ trap.

The PE induced by the addition of litter was calculated as:

$$
PE = (Rs \text{ soil with litter}) - (Rs \text{ control soil})
$$
 (3)

where Rs control soil is the $CO₂$ emitted by the control soil and Rs soil with litter is the $CO₂$ evolved from the soil treated with litter.

$$
P_{\text{litter}} = \frac{(R_{\text{t}} \times A_{\text{t}^{13}}) - (R_{\text{t}} \times A_{\text{s}^{13}})}{A_{\text{c}^{13}} - A_{\text{s}^{13}}}
$$
(4)

 P_{litter} is the proportion of soil CO_2 efflux derived from litter C. This equation was also used to partition the contributions of litter- and SOM-derived C to organic C recovered in the $K₂SO₄$ extracts from fumigated (F) and non-fumigated (NF) soils. The MBC derived from litter (and by difference from SOM) was calculated as in Paterson and Sim [\(2013](#page-9-0)):

$$
MBC_{\text{litter}} = \left[(P_{\text{litter}} \times C)_{\text{F}} - (P_{\text{litter}} \times C)_{\text{NF}} \right] / K_{\text{EC}} \tag{5}
$$

The relationship between the specific priming effect (PE) and incubation time was best explained and fitted by two different models (growth and peak functions) for the natural forest and coniferous plantation.

Statistics

A repeated ANOVA was performed to analyze the temporal effects of leaf-litter addition on SOM mineralization (released $CO₂$) and PE. A t test was used to test the effects of litter addition on the contents of DOC, MBC, PLFAs, etc. at a significant level of $p < 0.05$ (SPSS version 21, SPSS Inc., Chicago, IL, USA). Changes in the composition of main microbial groups were analyzed using principal component analysis (PCA; proportional mol% of PLFA) in PRIMER (v6.1.15, Primer-E Ltd., Plymouth UK). Analysis of similarities (ANOSIM) based on Bray-Curtis distances was used to test the differences in microbial communities between the litter addition and control treatments (Oksanen et al. [2007](#page-9-0)).

Results

Soil organic matter and litter mineralization

Leaf-litter addition increased the total $CO₂$ efflux (determined as cumulative CO_2 produced per gram SOC) by $44 \pm 3\%$ and $37 \pm 1\%$ in natural forest and plantation soil, respectively $(p < 0.05,$ Fig. 1). The total amount of SOM-mineralized C in the natural forest soil (20.5 g C kg⁻¹ SOC) was higher than that under plantation (16.2 g C kg⁻¹ SOC) over 195 days (Fig. 1).

Litter addition decreased the amount of soil-derived C in MBC $(qCO₂)$ of natural forest, but not for the plantation soil (Table [2\)](#page-4-0). Cumulative litter mineralization (normalized per gram microbial biomass C, i.e., $qCO₂$) in soils from the natural forest was higher (26%, $p < 0.01$) than under the plantation after 195 days. The natural forest and plantation had comparable CUE (similar $qCO₂$) within control soil, whereas natural forest had higher (up to 10%) soil CUE (lower qCO_2) than plantation within soil with litter addition (Table [2\)](#page-4-0).

Priming effect

In natural forest, the $CO₂$ production from SOM mineralization after litter addition increased by $13.6 \pm 0.04\%$ (i.e., positive PE), but decreased by $4.1 \pm 0.17\%$ in plantation soil (i.e., negative PE, Fig. [2\)](#page-4-0). Remarkably, a slow positive PE (1.1 μg $C-CO₂ g⁻¹ SOC)$ in natural forest soil was observed in the initial stage of ca. 50 days (Fig. [2\)](#page-4-0). However, in plantation soil, the cumulative PE (2.0 μg C-CO₂ g⁻¹ SOC) remained negative during the initial stage (Fig. [2](#page-4-0)). This demonstrated that during the initial intensive phase, litter decomposition exceeded the negative PE. After a period of 50 days, the direction of PE in plantation soil switched rapidly from negative

Fig. 1 Cumulative total $CO₂$ efflux during a 195-day incubation period in natural forest (NF) and plantation forest (PF) soils in subtropics. Results are means \pm standard errors ($n = 4$). SE is not presented if it is smaller than the symbol. Green arrows indicate the effect of litter addition

Table 2 Metabolic quotients (qCO_2 , means \pm standard errors) for soil-derived and litter-derived C (based on δ^{13} C). Letters indicate significant differences between the treatments (with or without litter addition) within each forest. Asterisk means significant difference between natural and plantation forests in litter addition treatment. Higher $qCO₂$ values indicate greater $CO₂$ release per microbial biomass unit (lower C use efficiency)

Treatments	Total $qCO2$	Soil $qCO2$	Litter $qCO2$
Natural forest			
Control	$2.63 \pm 0.01a$	$2.63 \pm 0.01a$	
Litter addition	$2.83 \pm 0.07a^*$	2.40 ± 0.06 b [*]	$7.87 \pm 0.19*$
Plantation forest			
Control	2.44 ± 0.10	$2.44 \pm 0.10a$	
Litter addition	$3.24 \pm 0.16a$	$2.64 \pm 0.11a$	6.24 ± 0.25

to positive. However, during the slow phase of litter decomposition, the PE under plantation increased drastically. The magnitude of PE in natural forest soil was similar to that of plantation soil in the later stage (i.e., after 50 days; Fig. 2). The cumulative positive PE in plantation soil during the later stage balanced out the negative PE in the initial stage, indicating similar effects on SOM storage after litter addition.

Microbial biomass

Litter addition increased MBC in natural forest soil, but not under plantation (Table [3\)](#page-5-0). The MBC under natural forest was

higher than that under plantation in soil with and without litter addition (Table [3\)](#page-5-0). In natural forest, litter addition increased soil-derived MBC compared to the control (by up to 31%, $p < 0.05$), whereas MBC decreased by 13% under plantation (Fig. [3,](#page-5-0) $p < 0.05$). The MBC derived from litter in natural forest soil (24 ± 4.6 mg C kg⁻¹ soil) was similar to that in plantation soil (27 ± 3.2 mg C kg⁻¹ soil; Fig. [3\)](#page-5-0). The percentage of litter-derived MBC to the total MBC in plantation was twice that in natural forest (Fig. [3\)](#page-5-0).

Effects of litter addition on composition of main soil microbial groups

In plantation soil, litter addition affected both microbial biomass (Fig. [3](#page-5-0)) and the composition of main microbial groups (Fig. [4\)](#page-5-0), with proportional increase of fungi and gram-positive bacteria compared to soil without litter. Litter addition increased ratio of gram-positive to gram-negative bacteria (G+/G−) and fungal to bacterial ratio (F/B) in the plantation soil. This result suggests changes occurred in microbial substrate utilization under plantation, but remained stable under the natural forest (Fig. [4](#page-5-0)). The main separation of the natural and plantation forest soils according to the PLFA composition along PC1 (explained by 62.8%) indicates that there were strong differences between microbial communities in two forest soils (Fig. [5\)](#page-6-0). The PCA separation of the forests between soils with and without litter addition based on the PLFA

Fig. 2 Priming effects after addition of 13 C labeled Chinese fir litter during a 195-day incubation period in two subtropical forest soils, explained by the main mechanism of microbial substrate adaptation. The priming dynamics were estimated by two models (growth and peak functions) for the natural and plantation forests, respectively. The peak

function for plantation forest that represents the shifts of priming effect from negative to positive. There are two stages, i.e., microbial substrate adaptation (initial stage) and rapid priming of SOM decomposition (later stage). Results are means \pm standard errors ($n = 4$). NS non-significant difference. * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$

Table 3 Dissolved organic C and microbial biomass C under natural evergreen broad-leaved forest and Chinese fir plantation as affected by leaf litter addition during a 195-day incubation period

Results are means \pm standard errors ($n = 4$). Letters following the means in the same row denote significant differences between control and litter addition treatments at $p < 0.05$

composition was much less expressed for the natural forest soil compared to the plantation soil (Fig. [5\)](#page-6-0).

Discussion

High specific C mineralization (expressed as $CO₂$ per gram SOC) indicates lower SOM recalcitrance (Mikan et al. [2002](#page-8-0); Fierer et al. [2003](#page-8-0); Salomé et al. [2010](#page-9-0)). Specific C mineralization in control soil under natural forest was up to 27% greater $(p < 0.05)$ than under plantation forest (Fig. [1](#page-3-0)). Therefore, SOM in natural forest is less recalcitrant than under plantation. Litter addition triggered up to 1.4 times higher mineralization of total $CO₂$ efflux than that in soil without litter addition, which caused an increase (up to 14%) in SOM decomposition under natural forest. In contrast, litter addition increased the total $CO₂$ efflux by 37% in plantation soil, whereas SOM decomposition slightly decreased by 4.1%. Consequently,

Fig. 3 Contribution of soil organic matter (SOM) and litter-derived C to total microbial biomass C (MBC) after a 195-day incubation period in natural and plantation forest soils. The red and blue arrows show the positive (+PE) and negative (−PE) priming effects in microbial biomass

litter addition triggered positive PE under natural forest but negative PE under plantation. This confirmed a high susceptibility of SOM to decomposition in the presence of litter under natural forest. Thus, our first hypothesis that litter addition would stimulate microbial-mediated decomposition of SOM in natural forest soils was confirmed.

Fig. 4 The ratios of gram-positive bacteria to gram-negative bacteria (G+/G− ratio) and fungi to bacteria (F/B ratio) after a 195-day incubation period in two subtropical forest soils. Results are means ± standard errors $(n = 4)$. *Significant difference between control and litter addition treatments at $p < 0.05$

Fig. 5 Principal component analysis (PCA) of the proportional mol% of the total microbial community of the natural forest (NF) and plantation forest (PF). The percentage of variance explained by each PC is shown in parentheses. The green arrow indicates the effect of litter addition

The differences in the direction of the PE depend on N availability, specifically exchangeable NH_4^+ (Chen et al. [2014;](#page-8-0) Li et al. [2017\)](#page-8-0). In general, positive PE owing to the "N mining" occurs when N availability is a limiting factor for microorganisms (Kuzyakov [2002](#page-8-0); Fontaine et al. [2003](#page-8-0); Dijkstra et al. [2013;](#page-8-0) Chen et al. [2014](#page-8-0)), whereas the "preferential substrate utilization" of litter instead of SOM induces negative PE (Kuzyakov [2002;](#page-8-0) Fontaine et al. [2004,](#page-8-0) [2011\)](#page-8-0). However, N availability in the natural broad-leaved forest soil is higher than that in plantation soil (Table [1\)](#page-2-0), so the positive PE is opposite to the proposed mechanisms.

Microorganisms in natural forest soil are usually adapted to utilize high-quality litter (low C/N, high N availability, diverse litter sources). These microorganisms need to mine more SOM to meet their demand for N when supplied with lowquality leaf litter. Microorganisms in natural forest mainly utilize litter-derived C as an energy source rather than for microbial growth to mineralize SOM, resulting in positive PE in natural forest (Schimel and Weintraub [2003](#page-9-0)). This can be supported by the higher litter-derived qCO_2 in natural for-est than that in plantation (Table [2](#page-4-0) and Fig. [3](#page-5-0)). Higher $qCO2$ in soil under natural forest indicates that microorganisms have

Fig. 6 Schematic diagram illustrating positive and negative priming effects caused by additions of 13C-labeled litter (from the plantation) into the soils of natural and plantation forests. The red lines show processes for microbial decomposition and assimulation of soil organic matter (SOM); the blue lines identify processes for added labeled leaf-litter. The thickness of the line indicates the magnitude of the process

lower C use efficiency (CUE) than that in plantation soil. The higher CUE (lower $qCO₂$) and more SOM-derived microbial biomass after litter addition relative to that of the control under natural forest promoted the growth of microorganisms better adapted to degrade SOM. Consequently, microorganisms in natural forest prefer to decompose SOM rather than added recalcitrant litter, thereby driving the positive PE (Creamer et al. [2015](#page-8-0)).

In contrast, microorganisms in plantation soil prefer to utilize the litter rather than SOM, as they have adapted over 42 years to Chinese fir litter and thus decompose them faster than that in natural forest soil (Veen et al. [2015](#page-9-0); Yu et al. [2015\)](#page-9-0). Microorganisms in plantation soil dominated by fungi, which favor utilization of litter-derived C (Guo et al. [2016](#page-8-0); Kramer et al. [2012](#page-8-0)) to meet their growth demands and thus incorporate more litter-derived C into microbial biomass (Fig. [3\)](#page-5-0). Litter addition decreased soil CUE (higher $qCO₂$) relative to the control soil under plantation, but microorganisms in plantation soil had higher litter CUE (lower $qCO₂$) than natural forest. Consequently, litter addition stimulated the growth of microorganisms less specialized to degrade SOM, resulting in negative PE. Thus, our second hypothesis was also confirmed.

Despite the differences in the direction and magnitude of priming between two soils immediately after litter addition, positive priming consistently increased in later stages (Fig. [2\)](#page-4-0). These differences in SOM decomposition after litter addition between the two soils are related to microbial community composition (Creamer et al. [2015](#page-8-0)). Soil priming reflects the activity of certain microbial groups (particularly fungi) more than others (De Graaff et al. [2010](#page-8-0); Fontaine et al. [2011](#page-8-0); Garcia-Pausas and Paterson [2011](#page-8-0); Paterson and Sim [2013](#page-9-0); Lü et al. [2015\)](#page-8-0). Priming in natural forest soil could not arise from changes in microbial community composition, as microbial community composition remained stable after litter addition (Figs. [4](#page-5-0) and [5](#page-6-0)). Instead, the high CUE and SOM-derived microbial biomass suggested that these microorganisms mainly K-strategists (Blagodatskaya et al. [2014\)](#page-8-0)—are favored by SOM utilization. The priming intensity in natural forest soil might be driven by microbial communities and their activities rather than community composition. However, the composition of main soil microbial groups changed after litter addition in plantation. Microbial community composition is associated with shifts in microbial substrate utilization (Collins et al. [2016\)](#page-8-0) and can underpin the PE direction (Blagodatskaya and Kuzyakov [2008;](#page-8-0) Dungait et al. [2011\)](#page-8-0). Litter addition increased the F/B ratio in the plantation soil, which implied changes in microbial substrate utilization (Shihan et al. [2017;](#page-9-0) Lin et al. [2018\)](#page-8-0), whereas there were no changes of F/ B ratio under natural forest. Consequently, during the intensive litter mineralization stage (initial stage ca. 50 days), negative PE in plantation soil is mainly attributed to the preferential substrate utilization (Blagodatskaya and Kuzyakov [2008](#page-8-0);

Garcia-Pausas and Paterson [2011](#page-8-0)). At later stages, the PE strongly exceeded litter decomposition and was accompanied by decreased incorporation of SOM-derived C into microbial biomass. This indicated that the PE were possible due to microbial shifts, i.e., from fast- to slow-growing (e.g., fungi) litter feeding populations. This also suggested a new mechanism of real PE, primarily from the litter-feeding microorganisms regulating PE (Miltner et al. [2009](#page-9-0), [2012;](#page-9-0) Shahbaz et al. [2017\)](#page-9-0).

Overall, the microbial community in natural forest soil continued to decompose SOM (positive PE) and produced more microbial biomass from SOM. The increased SOM-derived microbial populations accelerated SOM decomposition which depended on activity and biomass of microbial community rather than changes in the composition of main soil microbial groups. However, microorganisms in plantation soil switched from SOM mineralization to incorporation of litter C into microbial biomass, resulting in negative PE and soil C accumulation.

Conclusions

Two PE mechanisms (Fig. [6\)](#page-6-0) were proposed in soils under subtropical natural forest and 42-year-old coniferous plantation forest. The selective microbial decomposition of Chinese fir leaf-litter leads to positive PE in natural forest soil, but negative PE under plantation. This suggests that the input of Chinese fir litter following conversion of natural broad-leaved forests to plantations may accelerate SOM decomposition but would have subsequently been followed by negative priming with stand development due to microbial adaptation to plantation litter. Consequently, the priming-related soil C losses after subtropical forest conversion may ultimately have little or no long-term impact on soil C stocks. Therefore, the conversion of natural forests to plantations should consider divergent impacts of leaf-litter inputs on SOM priming. Further research efforts should focus on examining the effects of reciprocal litter transplant, i.e., home-field advantage of litter decomposition on PE under contrasting litter quality to assess how forest conversion with different tree species impacts on soil C dynamics.

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