




Article

Soil Microbial Diversity, Biomass, and Activity in Two Pine Plantations of Southern Italy Treated with Prescribed Burning

Elisabetta Giuditta ¹, Rossana Marzaioli ^{1,*}, Assunta Esposito ¹, Davide Ascoli ²,
Adriano Stinca ¹, Stefano Mazzoleni ³ and Flora A. Rutigliano ¹

¹ Department of Environmental Biological and Pharmaceutical Sciences and Technologies, University of Campania “Luigi Vanvitelli”, Via Vivaldi 43, 81100 Caserta, Italy; elisabetta.giuditta@gmail.com (E.G.); assunta.esposito@unicampania.it (A.E.); adriano.stinca@unicampania.it (A.S.); flora.a.rutigliano@unicampania.it (F.A.R.)

² Department of Agricultural, Forest and Food Sciences, University of Turin, Largo Paolo Braccini 2, 10095 Grugliasco (TO), Italy; d.ascoli@unito.it

³ Department of Agricultural Sciences, University of Naples Federico II, Via Università 100, 80055 Portici (NA), Italy; stefano.mazzoleni@unina.it

* Correspondence: rossana.marzaioli@unicampania.it

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Abstract: Microbial diversity plays a crucial role in ecosystem processes, including organic matter decomposition and nutrient cycling. This research explores the effect of prescribed burning (PB) on soil microbial diversity, as well as biomass and activity in Mediterranean pine plantations. In burned and adjacent unburned plots of *Pinus pinea* and *P. pinaster* plantations of Southern Italy protected areas, the fermentation layer and the 5 cm thick layer of mineral soil underneath were sampled at intervals during the first year after PB. The experimental protocol encompassed measurements of total microbial abundance (C_{mic} and soil DNA), fungal mycelium, fungal fraction of C_{mic} , microbial activity, bacterial genetic diversity (16S rDNA PCR-DGGE), microbial metabolic quotient (qCO_2), and C mineralization rate (CMR), as well as physical and chemical soil properties. PB caused only temporary (up to 3 h–32 d) reductions in C_{mic} , DNA amount, fungal mycelium, respiration, and CMR in the *P. pinaster* plantation, and had no appreciable negative effect on the microbial community in *P. pinea* plantation, where fire intensity was lower because of less abundant litter fuel. In either plantation, PB did not generally reduce bacterial genetic diversity (evaluated as band richness, Shannon index, and evenness), thus, also accounting for the fast recovery in microbial growth and activity after high-intensity PB in *P. pinaster* plantation. While confirming PB as a sustainable practice to reduce wildfire risk, also supported by data on plant community obtained in the same plantations, the results suggest that an integrated analysis of microbial diversity, growth, and activity is essential for an accurate description of PB effects on soil microbial communities.

Keywords: *Pinus pinea* plantation; *Pinus pinaster* plantation; prescribed burning; microbial biomass; microbial activity; bacterial genetic diversity

1. Introduction

Soil microbial community plays a key role in terrestrial ecosystems, by being involved in several ecosystem services. Indeed, it provides supporting services (soil formation, nutrient cycling, and plant growth), as well as regulating services (i.e., climate and gas regulation, C sequestration, water purification, disease and pest regulation, and bioremediation) and provisioning ones (i.e., the supply of food, fiber, fuel, genetic resources, chemicals, and pharmaceuticals) [1–3].

Soil microorganisms generally respond to changes in environmental conditions much faster than physical and chemical properties such as soil organic C or total N content [4–6]. Changes in soil microbial community, therefore, are useful indicators of the effects of stress or disturbance factors on soil. Research on soil microbial community currently concerns the evaluation of changes in quantity or biomass, structure (ranging from community fingerprint to species identification), and activity as indicators of environmental changes [7] by using traditional biochemical and microscopy techniques, as well as the newer molecular procedures [8]. However, only a few studies have simultaneously used microbial indicators of quantity or biomass, structure, and activity, notwithstanding the relationship among them is still unclear; therefore, studies in this research field should include the assessment of each of these three groups of indicators [7].

Common sources of disturbance in the Mediterranean ecosystems are wildfires, which affect a substantial amount of the forest cover, especially in concomitance with summer drought [9,10]. By causing a general reduction in the forest understory and tree crown cover and changes in physical, chemical, and biological properties of soil, wildfires can adversely affect soil functions. Indeed, soil alteration induced by fire could favour erosive processes and nutrient loss, thus affecting nutrient cycling [11], microbial activity, and CO₂ emission from soil [12,13]. In order to reduce the fire risk in Mediterranean forests, the prescribed burning (hereafter PB) technique has been increasingly employed in the last years [14,15]. PB is applied under specific operative conditions (known as “prescriptions”) with the main goal of reducing the amount of surface dead fine fuels for fire hazard abatement, while minimizing undesired fire effects on ecosystems, including the soil component [14,16].

Several studies have shown that PB application is a useful fuel management tool that reduces wildfire hazard and severity, thus, improving ecosystem resilience to wildfire, particularly in Mediterranean pine plantations [14,16–22]. However, current understanding of the effects of prescribed burning on ecosystem components and processes is incomplete. Whereas extensive literature is available on the PB effects on vegetation and plant physiology [14,23–29], the knowledge of the effects of PB on soil components and processes is still unsatisfactory and mainly concerns soil chemical properties, while biological properties have not been widely investigated, until now [30]. Moreover, the biological data available are not fully consistent because both negative, positive, or neutral effects of PB have been reported depending on the soil initial characteristics, vegetation, fire intensity and frequency [30–33]. In addition, research on the effects of PB on soil microbial community has been mostly focused on total microbial biomass and activity [34–36], whereas effects on all variables linked to soil microbial community (quantity/biomass, structure, and activity) after PB treatment have been investigated only occasionally, i.e., in Texas grassland [37], Oklahoma oak forest [33], and British Columbia fir forest [38]. Moreover, the soil microbial diversity is not well understood in Mediterranean areas treated with PB.

With the aim to provide useful information on the sustainability of this practice for forest management in Mediterranean ecosystems, we investigated soil microbial diversity (as bacterial genetic diversity), abundance (as biomass and DNA amount), and activity (as respiration and indexes of microbial metabolism), and the relationships among these variables, in two pine plantations of Southern Italy treated with PB. We chose two plantations (dominated respectively, by *P. pinea* and *P. pinaster*) that differed for litter amount in order to obtain different fire intensity.

We hypothesize the following: (1) prescribed burning has no or slight effect on microbial biomass, activity, and diversity of fermentation layer and underlying 5 cm soil in both plantations and (2) the change in microbial diversity is a useful indicator of the effect of PB on soil microbial community that has to be associated to indicators more commonly investigated, i.e., variations in microbial biomass and activity.

2. Materials and Methods

2.1. Study Areas

The study was carried out in two pine plantations dominated by *P. pinea* L. and *P. pinaster* Aiton subsp. *pinaster*, respectively (Figure 1). The *P. pinea* plantation (40°57'39.79'' N and 13°59'50.09'' E) is located in the Castel Volturno Nature Reserve [39], a sandy coastal area located between sea level and 9 m of altitude. The mean age of trees was 50 years [24]. The understory mainly consists of flammable plants, such as *Phillyrea angustifolia* L., *Rhamnus alaternus* L. subsp. *Alaternus*, and *Asparagus acutifolius* L. [24,40]. The climate is typically Mediterranean, with hot dry summers and cool wet winters. Data from the meteorological station of Ischitella for 1974 to 2012 report a mean annual temperature of 13.6 °C and a mean annual rainfall of 761.3 mm [24]. Soil is a *Calcacoric Arenosol* according to the FAO system of soil classification [41].

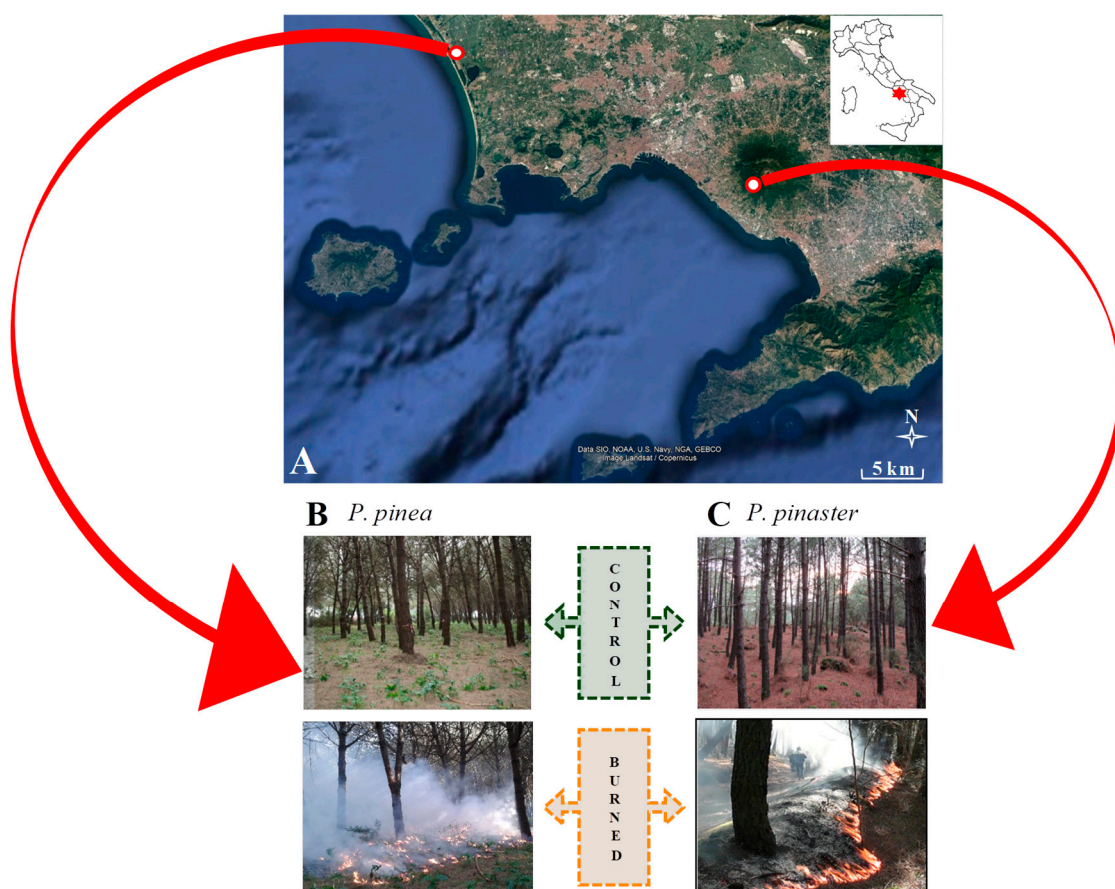


Figure 1. Study areas in the Campania region, Southern Italy, Google Earth Pro, July 4, 2019 (A). Control and burned plots in *P. pinea* (B) and *P. pinaster* (C) plantations.

The *P. pinaster* plantation (40°48'42.84'' N and 14°24'46.86'' E) is located in the Tirone Alto Vesuvio Nature Reserve, within the Vesuvius National Park, at about 600 m a.s.l. and 20% slope. The mean age of trees was 37 years [42]. The understory consists of *Robinia pseudoacacia* L., *Genista etnensis* (Raf.) DC., *Cytisus scoparius* (L.) Link subsp. *scoparius*, and *Quercus ilex* L. subsp. *ilex*. The mean annual temperature in the interval 1951–2014 was 16.1 °C and the mean annual precipitation was 766 mm [43]. Soil is a *Lepti-Vitric Andosol (Skeletal)* according to the FAO system of soil classification [41].

Overall, the *P. pinea* plantation showed a higher number of plant species in the forest understory (9.5 ± 0.7) as compared with *P. pinaster* (6.91 ± 2.5) plantation, as evaluated, respectively, on 50 m² plots (Esposito et al., unpublished results) and 78.5 m² plots (Stinca et al., unpublished results). It

also showed lower tree density and higher values of shrub and herb cover, as well as herb height as compared with *P. pinaster* (Table 1).

Table 1. Stand structure, fuel load, and operative conditions during prescribed burning in two pine plantations ¹.

	<i>P. pinea</i>	<i>P. pinaster</i>
Stand Structure		
Tree density (n. ha ⁻¹)	954 (±178)	1294 (±243)
Tree cover (%)	65 (±7.1)	62.3 (±5.9)
Tree high (m)	13.5 (±2.1)	11.7 (±0.7)
Shrub cover (%)	70.0 (±14.1)	37.1 (±24.6)
Shrub high (m)	1.8 (±3.5)	1.9 (±1.2)
Herb cover (%)	62.5 (±12.6)	2.1 (±1.7)
Herb high (cm)	32.5 (±5.0)	15.0 (±0.7)
Fuel load (t ha⁻¹)		
Litter (<6 mm)	6.4 (±1.1)	14.8 (±1.7)
Woody litter (6–25 mm)	1.4 (±1.0)	2.7 (±2.3)
Herb fuels	0.54 (±0.21)	Not analysed
Weather data ²		
Air temperature (°C)	18 (±0.65)	16 (±0.57)
Relative humidity (%)	54 (±4.1)	52 (±3.7)
Litter moisture (%)	32 (±3.8)	38 (±4.1)
Wind speed (km h ⁻¹)	2.7 (±0.9)	4.2 (±1.8)
Number of days since rain before the treatment	8	17
Fire behavior		
Ignition pattern	Backfire	Backfire
Flame length (m)	<0.5	<1
Rate of spread (m min ⁻¹)	0.14 (±0.04)	0.19 (±0.03)
Fireline intensity (kW m ⁻¹)	<50	<150
Litter mean maximum temperature (°C)	347 (±202)	574 (±205)
F-layer mean maximum temperature (°C)	95 (±136)	438 (±97)
Mean residence of temperature (s) above 100 °C in the litter	139 (±90)	230 (±42)
Mean residence of temperature (s) above 100 °C under F-layer	30 (±67)	180 (±70)

¹ Data are means ± standard deviations, except for number of days since rain, flame length, and fireline intensity and ² weather conditions were monitored, each 30 min during the burn, by a portable weather station (Kestrel 4500) positioned at 2 m above the ground.

2.2. Prescribed Burning Treatment

In both plantations, the PB was applied in March 2014 with the main aim to reduce the fire hazard by reducing surface fuel loadings and interrupting fuel continuity. Either study area was divided into the following two plots about 0.50 ha each: treated with PB (burned) and untreated (control). The operative conditions during the treatment in terms of weather conditions, fuel load (as dry weight), and fire behaviour are reported in Table 1. The arrival time of the fire was assessed by using 9 K-type thermocouples (0.4 mm in diameter) positioned in the litter and 9 just below the fermentation layer in *P. pinea* plantation, and 7 thermocouples in the litter and 6 under the fermentation layer in *P. pinaster* plantation. From each thermocouple we extracted maximum temperature and the residence time (s) above 100 °C (Table 1).

PB was more intense in *P. pinaster* than in the *P. pinea* plantation (<150 kW m⁻¹ vs. <50 kW m⁻¹), because of more abundant fine litter fuel (14.8 ± 1.7 vs. 6.4 ± 1.1 t ha⁻¹). This resulted in higher temperatures and more prolonged residence times above 100 °C in the litter and F-layer (Table 1). In both plantations the litter layer thickness had not yet recovered to the prefire levels, at least up to 18 months after the treatment [44].

2.3. Sampling Protocol and Laboratory Measurements

Samples of the fermentation layer (F-layer) and underlying 5 cm soil (S-layer) were collected in 6 subplots (40 × 40 cm) of burned and unburned plots. In each subplot, all fermentation layer was collected; successively, five cores of S-layer were collected by a cylindrical sampler (diameter 6 cm and height 5 cm) and then mixed to obtain a homogeneous sample. The sampling was performed at 3 h, and 33, 89, 189, and 363 days after PB in the *P. pinea* plantation and at 3 h, and 32, 87, 182, and 371 days after PB in the *P. pinaster* plantation.

Limited to the first sampling (3 h), the weight of fermentation layer (after drying at 75 °C) was determined.

At each sampling time, the F-layer and S-layer (the latter sieved at 2 mm mesh) were analyzed for water content, total and extractable organic C, total microbial biomass, DNA amount, fungal mycelium, microbial activity, and bacterial genetic diversity. In addition, pH, total and mineral N (as ammonium and nitrate) were determined for S-layer samples.

Water content of F- and S-layer samples was measured by gravimetric method [45]. S-layer pH was determined on a water suspension of air-dried soil (1:2.5 ratio) using a calibrated electrode (Hanna Instruments mod. HI1230) [46]. Total N was determined on dried (75 °C) and pulverized soil with a NCS Elemental Analyzer (Thermo FlashEA 1112). Ammoniacal and nitric N contents were determined on fresh soil stored at 4 °C until measurement. Mineral nitrogen was extracted from soil with 0.5 M K₂SO₄ (1:5 soil: extractant) and measured by selective electrodes for NH₄⁺-N (ORION, Mod. 9512BNWP) and NO₃⁻-N (ORION, Mod. 9707BNWP) [47]. Total organic carbon (C_{org}) of dried F- and S-layer samples was measured by humid digestion in 0.33 M K₂Cr₂O₇ [48,49].

Fresh F-layer and S-layer, stored at 4 °C until measurement, were used to determine total microbial biomass C (C_{mic}), fungal mycelium amount, and respiration activity. Moreover, the following two indexes of microbial metabolism were calculated: metabolic quotient (qCO₂: g CO₂-C kg⁻¹ C_{mic} h⁻¹) [50], indicating the activity level of microbial community, and C mineralization rate (CMR: g CO₂-C kg⁻¹ C_{org} h⁻¹), representing the fraction of organic C mineralized in the time unit. In addition, fungal fraction of microbial C (C_{fung} % C_{mic}) was evaluated.

Total microbial biomass was assessed with the chloroform-fumigation extraction method [51]. Organic carbon, extracted with 0.5 M K₂SO₄ from chloroform-fumigated samples and not fumigated samples, was measured by chemical digestion with 0.066 M K₂Cr₂O₇. On the basis of the organic carbon (C_{org}) content in extracts of fumigated and non-fumigated samples, microbial biomass carbon (C_{mic}) was calculated [51]. This measure also provided data on the extractable organic C (C_{ext}) that was obtained from non-fumigated samples. The abundance of soil microorganisms was also deduced from total DNA, considered a robust indicator of microbial biomass [52]. DNA was extracted from 0.25 g of F- or S-layer samples (stored at -20 °C until use) with the FastDNA SPIN Kit for soil (Bio 101 Inc). Two subplots (3rd and 4th) of burned and unburned plots were used for assessing DNA content and bacterial diversity (see below) in four over five sampling times (3 h, and 33, 189, and 363 days in *P. pinea* plantation and 3 h, and 32, 182, and 371 days in *P. pinaster* plantation).

Total fungal mycelium was determined with the membrane filter [53]. The length of hyphae was determined with the intersection method [54]. The mass of mycelia was evaluated from the average values of cross section (9.3 μm²), density (1.1 g mL⁻¹), and dry mass of the hyphae (15% of the wet mass) [55]. From fungal mycelium and C_{mic} data, the fungal fraction of microbial C (C_{fung} % C_{mic}) was calculated [56].

The microbial activity was assessed as potential respiration, by measuring the CO₂ evolved from samples by gas chromatography (TRACE™ Ultra Gas Chromatograph) [57], modified. Samples were incubated in standard conditions (25 °C, 55% water-holding capacity, in the dark) for 1 h before analysis [58].

The metabolic quotient (qCO₂: g CO₂-C kg⁻¹ C_{mic} h⁻¹) and C mineralization rate (CMR: g CO₂-C kg⁻¹ C_{org} h⁻¹) were calculated from respiration, microbial biomass (C_{mic}), and total organic C (C_{org}) data.

The bacterial genetic diversity was evaluated by 16S rDNA PCR-DGGE on DNA extracts [59]. PCR amplification was performed using the primer set GC 968f/UNI 1401r [60,61] in the conditions described by Agnelli et al. [62]. PCR products were quantified on 1% agarose gel in 1X TBE buffer using Low DNA Mass Ladder 100 bp as a marker. Denaturing-gradient gel electrophoresis (DGGE) was performed using the DcodeSystem (Universal Mutation Detection System, Biorad). Operative conditions were 6% polyacrylamide gel and 35% to 50% denaturant gradient of urea and formamide (100% denaturant contained 7 M urea and 40% formamide). The gel was run at 60 °C under 200 V for 4 h. DGGE profiles of the bacterial community were analyzed with ImageJ ver.1.33, in order to obtain band number and band intensity for each lane [59]. The bacterial diversity was assessed in terms of richness (band number), Shannon index (H) and evenness index. The Shannon index was calculated by the equation, $H = -\sum(n_i/N) \times \ln(n_i/N)$, where n_i is the intensity of each band and N the sum of the intensities of all bands in the same lane [63]. The evenness index was calculated as the ratio between Shannon index and natural logarithm of richness, the latter corresponding to the maximum value of Shannon index for the observed band number.

2.4. Statistical Analysis

A normality test (Kolmogorov–Smirnov) was applied to datasets before parametric tests; the data were transformed by log10 when not normally distributed [64]. For each variable, two-way ANOVA (followed, if required, by Bonferroni test) was applied using treatment (burned or control) and sampling time (T) as factors. The significance level of the Bonferroni test, for burned vs. control comparisons was $p \leq 0.01$, according to Bonferroni method [64]. Moreover, the significance of differences between two considered plantations, for each variable, was assayed by *t*-test.

Correlations among parameters were assayed by Pearson's coefficient ($n = 60$ throughout except genetic data, for which $n = 16$).

All statistical analyses were carried out using the software SigmaPlot12.

3. Results

In both plantations, PB consumed litter but not the F-layer. In the *P. pinaster* plantation, an increase in the mass of the F-layer was observed ($839 \pm 373 \text{ g m}^{-2}$ before PB vs. $1231 \pm 331 \text{ g m}^{-2}$ after PB), suggesting that partially mineralized litter was transferred to this layer. The same did not occur in *P. pinea* plantation ($1140 \pm 277 \text{ g m}^{-2}$ before PB vs. $1135 \pm 367 \text{ g m}^{-2}$ after PB), indicating that litter had been entirely mineralized.

PB generally did not affect soil chemical characteristics in *P. pinea* plantation, however, a significant increase was found on the overall study period in N_{tot} , $\text{NH}_4^+\text{-N}$, and $\text{NO}_3^-\text{-N}$ concentrations (Table 2 (A)), and, limited to 89 and 363 days, in extractable C (C_{ext}) concentration of the F layer (Table 3 (A)). In this plantation, positive correlations were found between $\text{NH}_4^+\text{-N}$ concentration and soil DNA and fungal mycelium (respectively, $r = 0.50$ and $r = 0.68$, $p < 0.05$) and between $\text{NO}_3^-\text{-N}$ concentration and soil respiration, $q\text{CO}_2$, and CMR (respectively, $r = 0.45$, $r = 0.29$, $r = 0.463$, $p < 0.05$).

In *P. pinaster* plantation, burning treatment produced a slightly but significant increase in soil pH, and a significant decrease in $\text{NH}_4^+\text{-N}$ content, limited to 32 d and 182 d (Table 2 (B)), water content in both the F- and S-layers (Table 3 (B)). In either plantation, PB treatment did not affect the organic C concentration in F- or S-layer (C_{org} , Table 3).

Microbial variables also showed different responses to PB in the two plantations. A comparison of burned and control plots, showed that no negative effect was generally found in *P. pinea* plantation (Figure 2); however, a significant decrease in fungal mycelium was found in the S-layer, 3 h after the treatment (Figure 2F). The observed changes in respiration and $q\text{CO}_2$ (Figure 2I,K) were not clearly referable to fire treatment because they were sometimes higher in the control and sometimes the reverse. In *P. pinaster* plantation a negative effect of PB on some microbial variables was observed in both the F- and S-layers (Figure 3), but this effect was temporary. In particular, both layers showed a significant decrease in total C microbial biomass (C_{mic}) and fungal mycelium immediately after burning (3 h)

and generally up to 32 days thereafter (Figure 3A,B,E,F). A decrease in respiration (Figure 3I) and C mineralization rate (CMR, Figure 3M) was observed in F-layer at 3 h and 371 days after burning. The burned plots also showed a slight reduction in S-layer DNA content (Figure 3D). Unlike other microbial variables, the fungal fraction of C_{mic} and qCO_2 of the S-layer were higher in burned plots than in controls up to 32 d after treatment (Figure 3H,L).

Table 2. Mean (\pm standard deviation) values of soil pH, total (N_{tot}), and mineral N content (NH_4^+ -N and NO_3^- -N content) in the control and burned plots of *P. pinea* (A) and *P. pinaster* (B) plantations at different times after burning. Results of two-way ANOVA for burning and sampling time were reported at the bottom of A and B, respectively. N.S., nonsignificant. For each of 5 sampling times, significant differences between treatments (determined by Bonferroni test at $p \leq 0.01$) are indicated by different letters (a and b) in superscripts.

(A) Time after Burning (Sampling Date)	pH	N_{tot} (g kg ⁻¹ d.w.)	NH_4^+ -N (mg kg ⁻¹ d.w.)	NO_3^- -N (mg kg ⁻¹ d.w.)
3 h (12 March 2014)				
Control	7.2 (\pm 0.6)	2.0 (\pm 0.5)	1.6 (\pm 1.1)	4.6 (\pm 1.3)
Burned	6.6 (\pm 0.8)	2.6 (\pm 0.7)	4.1 (\pm 2.8)	7.0 (\pm 2.1)
33 d (14 April 2014)				
Control	6.8 (\pm 0.6)	2.2 (\pm 0.8)	11.8 (\pm 2.0)	6.2 (\pm 2.5)
Burned	6.4 (\pm 0.5)	3.1 (\pm 0.9)	13.7 (\pm 2.6)	5.8 (\pm 1.8)
89 d (9 June 2014)				
Control	6.5 (\pm 0.7)	2.7 (\pm 0.9)	7.6 (\pm 0.7)	7.6 (\pm 3.6) ^a
Burned	6.8 (\pm 0.5)	2.9 (\pm 0.7)	9.4 (\pm 3.9)	48.3 (\pm 23.4) ^b
189 d (17 September 2014)				
Control	6.7 (\pm 0.4)	2.1 (\pm 0.3)	0.9 (\pm 0.3)	24.0 (\pm 7.7) ^a
Burned	6.5 (\pm 0.5)	2.6 (\pm 0.7)	1.2 (\pm 0.4)	54.3 (\pm 26.2) ^b
363 d (10 March 2015)				
Control	6.7 (\pm 0.4)	2.2 (\pm 0.6)	13.7 (\pm 2.2)	2.4 (\pm 0.5)
Burned	6.5 (\pm 0.5)	2.4 (\pm 0.6)	15.9 (\pm 2.3)	2.4 (\pm 0.9)
Burning	N.S.	$p < 0.05$	$p = 0.001$	$p < 0.001$
Sampling time	N.S.	N.S.	$p < 0.001$	$p < 0.001$
(B) Time after Burning (Sampling Date)	pH	N_{tot} (g kg ⁻¹ d.w.)	NH_4^+ -N (mg kg ⁻¹ d.w.)	NO_3^- -N (mg kg ⁻¹ d.w.)
3 h (21 March 2014)				
Control	5.6 (\pm 0.5)	4.5 (\pm 3.0)	1.3 (\pm 1.0)	1.8 (\pm 0.3)
Burned	6.2 (\pm 0.4)	4.9 (\pm 4.8)	2.2 (\pm 1.5)	1.7 (\pm 0.1)
32 d (22 April 2014)				
Control	6.2 (\pm 0.5)	4.9 (\pm 3.2)	31.5 (\pm 15.6) ^a	4.0 (\pm 1.1)
Burned	6.4 (\pm 0.2)	1.2 (\pm 0.4)	14.3 (\pm 5.0) ^b	4.1 (\pm 0.4)
87 d (16 June 2014)				
Control	6.1 (\pm 0.5)	1.5 (\pm 0.9)	2.0 (\pm 0.2)	1.2 (\pm 0.2)
Burned	6.3 (\pm 0.5)	2.5 (\pm 1.8)	2.1 (\pm 0.2)	1.1 (\pm 0.1)
182 d (19 September 2014)				
Control	6.0 (\pm 0.5)	7.4 (\pm 3.3)	3.4 (\pm 1.6) ^a	2.9 (\pm 0.7)
Burned	6.2 (\pm 0.3)	7.5 (\pm 4.0)	1.7 (\pm 0.5) ^b	2.4 (\pm 0.5)
371 d (27 March 2015)				
Control	6.0 (\pm 0.5)	7.4 (\pm 3.6)	32.3 (\pm 8.7)	3.4 (\pm 1.0)
Burned	6.2 (\pm 0.3)	7.4 (\pm 2.9)	28.4 (\pm 12.3)	2.6 (\pm 1.0)
Burning	$p < 0.05$	N.S.	N.S.	N.S.
Sampling time	N.S.	$p < 0.001$	$p < 0.001$	$p < 0.001$

Table 3. Mean (\pm standard deviation) values of water content (WC), organic carbon (C_{org}), and extractable organic carbon (C_{ext}) content in the fermentation layer (F-layer) and in the 5 cm soil layer underneath (S-layer) in control and burned plots of *P. pinea* (A) and *P. pinaster* (B) plantations at different times after burning. Results of two-way ANOVA for burning and sampling time were reported at the bottom of A and B, respectively. N.S., nonsignificant. For each of 5 sampling times, significant differences between treatments (determined by Bonferroni test at $p \leq 0.01$) are indicated by different letters (a and b) in superscripts.

(A) Time after Burning (Sampling Date)	F-Layer			S-Layer		
	WC (%)	C_{org} (g kg ⁻¹ d.w.)	C_{ext} (g kg ⁻¹ d.w.)	WC (%)	C_{org} (g kg ⁻¹ d.w.)	C_{ext} (g kg ⁻¹ d.w.)
3 h (12 March 2014)						
Control	164 (± 44)	505.8 (± 46.9)	2.6 (± 0.7)	25 (± 7)	46.5 (± 4.6)	0.2 (± 0.1)
Burned	182 (± 20)	503.3 (± 42.8)	2.6 (± 0.4)	31 (± 8)	48.8 (± 11.1)	0.3 (± 0.04)
33 d (14 April 2014)						
Control	180 (± 27)	482.5 (± 27.6)	2.7 (± 0.3)	22 (± 7)	48.9 (± 16.4)	0.2 (± 0.03)
Burned	168 (± 9)	481.8 (± 51.5)	2.5 (± 0.4)	25 (± 14)	51.9 (± 12.3)	0.2 (± 0.1)
89 d (9 June 2014)						
Control	34 (± 10)	422.2 (± 60.3)	1.1 (± 0.3)	15 (± 11)	45.1 (± 5.4)	0.1 (± 0.04)
Burned	35 (± 11)	367.7 (± 37.1)	1.9 (± 0.3)	18 (± 14)	46.1 (± 7.7)	0.2 (± 0.04)
189 d (17 September 2014)						
Control	182 (± 81)	441.9 (± 30.7)	2.4 (± 0.9)	14 (± 3)	52.4 (± 7.4)	0.2 (± 0.1)
Burned	159 (± 38)	451.9 (± 50.3)	2.2 (± 0.5)	15 (± 3)	53.2 (± 12.7)	0.2 (± 0.04)
363 d (10 March 2015)						
Control	172 (± 21)	359.7 (± 50.2)	2.8 (± 0.7) ^a	29 (± 13)	48.5 (± 9.5)	0.3 (± 0.02)
Burned	154 (± 41)	337.6 (± 48.6)	4.4 (± 1.0) ^b	28 (± 11)	49.6 (± 9.5)	0.3 (± 0.1)
Burning	N.S.	N.S.	$p < 0.05$	N.S.	N.S.	N.S.
Sampling time	$p < 0.001$	$p < 0.001$	$p < 0.001$	$p < 0.001$	N.S.	$p < 0.001$
(B) Time after Burning (Sampling Date)	F-layer			S-layer		
	WC (%)	C_{org} (g kg ⁻¹ d.w.)	C_{ext} (g kg ⁻¹ d.w.)	WC (%)	C_{org} (g kg ⁻¹ d.w.)	C_{ext} (g kg ⁻¹ d.w.)
3 h (21 March 2014)						
Control	267 (± 92) ^a	413.9 (± 39.1)	2.6 (± 0.7)	54 (± 14)	82.8 (± 5.9)	0.3 (± 0.1)
Burned	147 (± 41) ^b	404.0 (± 36.5)	2.7 (± 1.0)	31 (± 13)	78.0 (± 4.4)	0.2 (± 0.1)
32 d (22 April 2014)						
Control	258 (± 76) ^a	426.6 (± 45.4)	4.3 (± 1.4)	58 (± 15) ^a	79.0 (± 8.2)	0.3 (± 0.1)
Burned	179 (± 33) ^b	421.1 (± 36.1)	2.9 (± 0.8)	21 (± 6) ^b	75.7 (± 15.3)	0.2 (± 0.1)
87 d (16 June 2014)						
Control	135 (± 40)	407.4 (± 35.7)	2.5 (± 0.8)	19 (± 6)	78.5 (± 11.7)	0.1 (± 0.1)
Burned	97 (± 34)	400.0 (± 53.3)	1.9 (± 0.3)	23 (± 9)	84.9 (± 14.9)	0.2 (± 0.03)
182 d (19 September 2014)						
Control	292 (± 32)	430.3 (± 46.3)	4.7 (± 1.0)	162 (± 32)	75.8 (± 12.7)	0.5 (± 0.2)
Burned	264 (± 41)	429.3 (± 47.1)	4.4 (± 1.1)	140 (± 47)	77.2 (± 4.6)	0.5 (± 0.2)
371 d (27 March 2015)						
Control	247 (± 55)	426.9 (± 36.3)	5.5 (± 1.3)	172 (± 80)	79.6 (± 12.8)	0.5 (± 0.1)
Burned	256 (± 49)	426.6 (± 48.7)	4.3 (± 0.9)	109 (± 86)	78.3 (± 9.1)	0.3 (± 0.1)
Burning	$p < 0.001$	N.S.	$p < 0.05$	$p < 0.001$	N.S.	$p < 0.05$
Sampling time	$p < 0.001$	N.S.	$p < 0.001$	$p < 0.001$	N.S.	$p < 0.001$

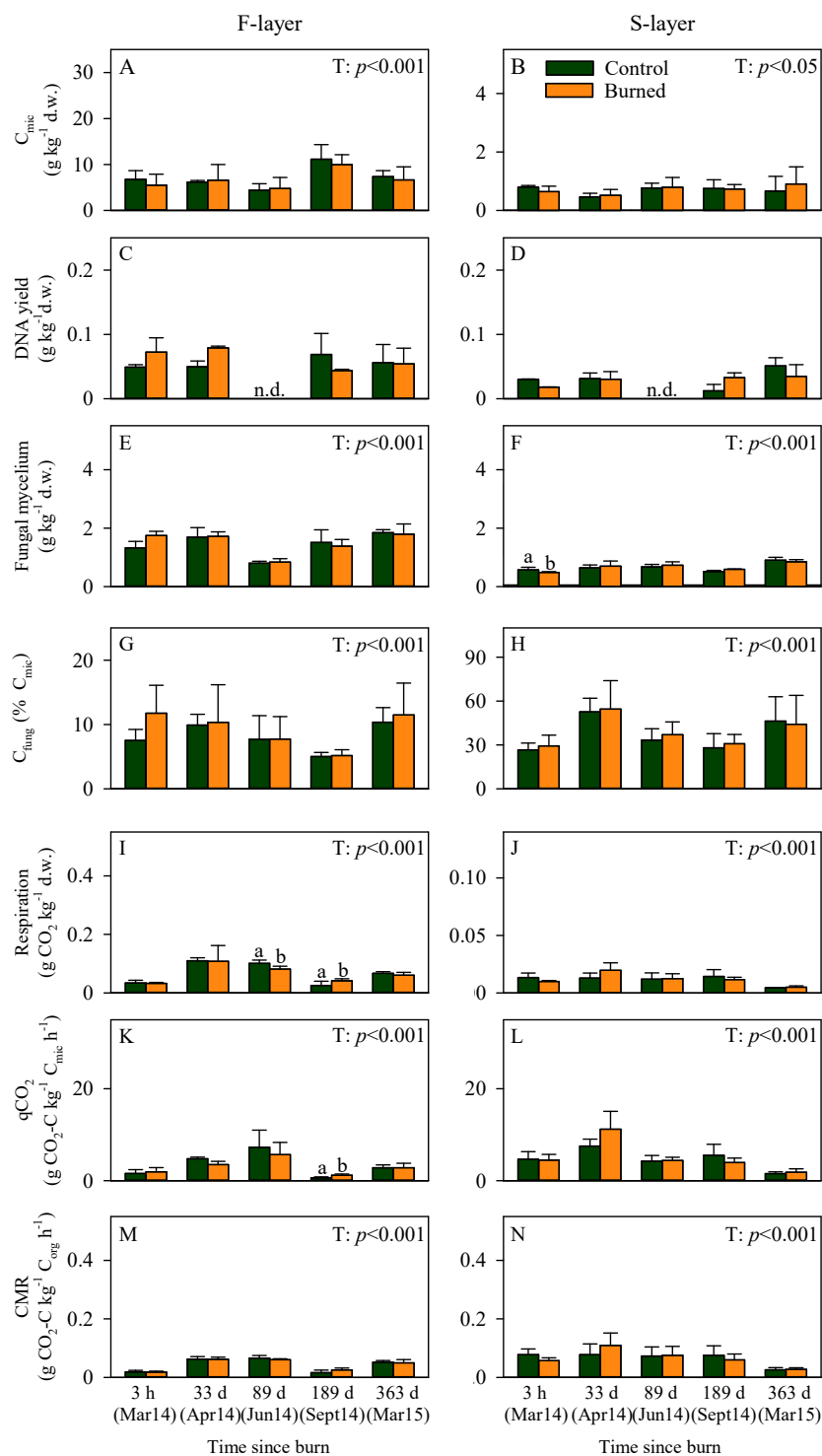


Figure 2. *P. pinea* plantation. Mean (+ standard deviation) values of microbial biomass C (C_{mic}) (A,B), DNA yield (C,D), fungal mycelium (E,F), fungal fraction of C_{mic} ($C_{fung} \% C_{mic}$) (G,H), respiration (I,J), metabolic quotient (qCO_2) (K,L), and carbon mineralization rate (CMR) (M,N) in the fermentation layer (F-layer), and in the 5 cm soil underneath (S-layer) of control and burned plots at different times (3 h, and 33, 89, 189, 363 d) after treatment. Each graph reports significant effects of burning (B) and sampling time (T) as assayed by two-way ANOVA (on the top), and of burning alone at each sampling time, by Bonferroni test (with different letters on bars, $p \leq 0.01$).

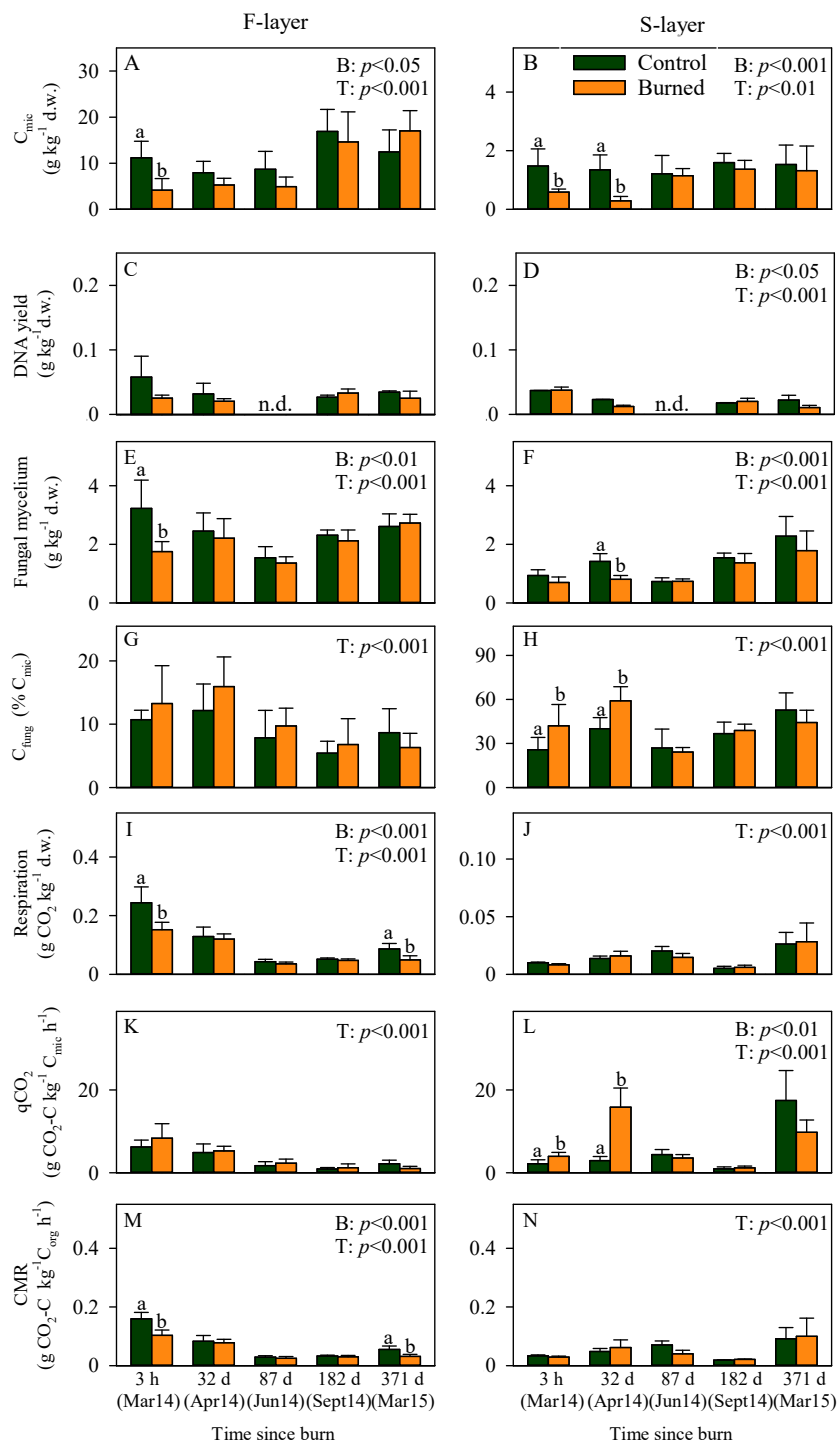


Figure 3. *P. pinaster* plantation. Mean (+ standard deviation) values of microbial biomass C (C_{mic}) (A,B), DNA yield (C,D), fungal mycelium (E,F), fungal fraction of C_{mic} (C_{fung} % C_{mic}) (G,H), respiration (I,J), metabolic quotient (qCO_2) (K,L) and carbon mineralization rate (CMR) (M,N) in the fermentation layer (F-layer) and in 5 cm soil layer underneath (S-layer) of control and burned plots at different times (3 h, and 32, 87, 182, 371 d) after the treatment. Each graph reports significant effects of burning (B) and sampling time (T) as assayed by two-way ANOVA (on the top), and of burning alone at each sampling time, by Bonferroni test (with different letters on bars; $p \leq 0.01$).

From the analysis of the DGGE profiles or bacterial genetic fingerprints of the F-layer (Figure 4A,C) and S-layer (Figure 4B,D) of considered plantations, no significant differences were generally observed

for bacterial richness, Shannon index, and evenness between burned and unburned plots (Table 4), except for richness in *P. pinea* S-layer, which was slightly lower in burned plot.

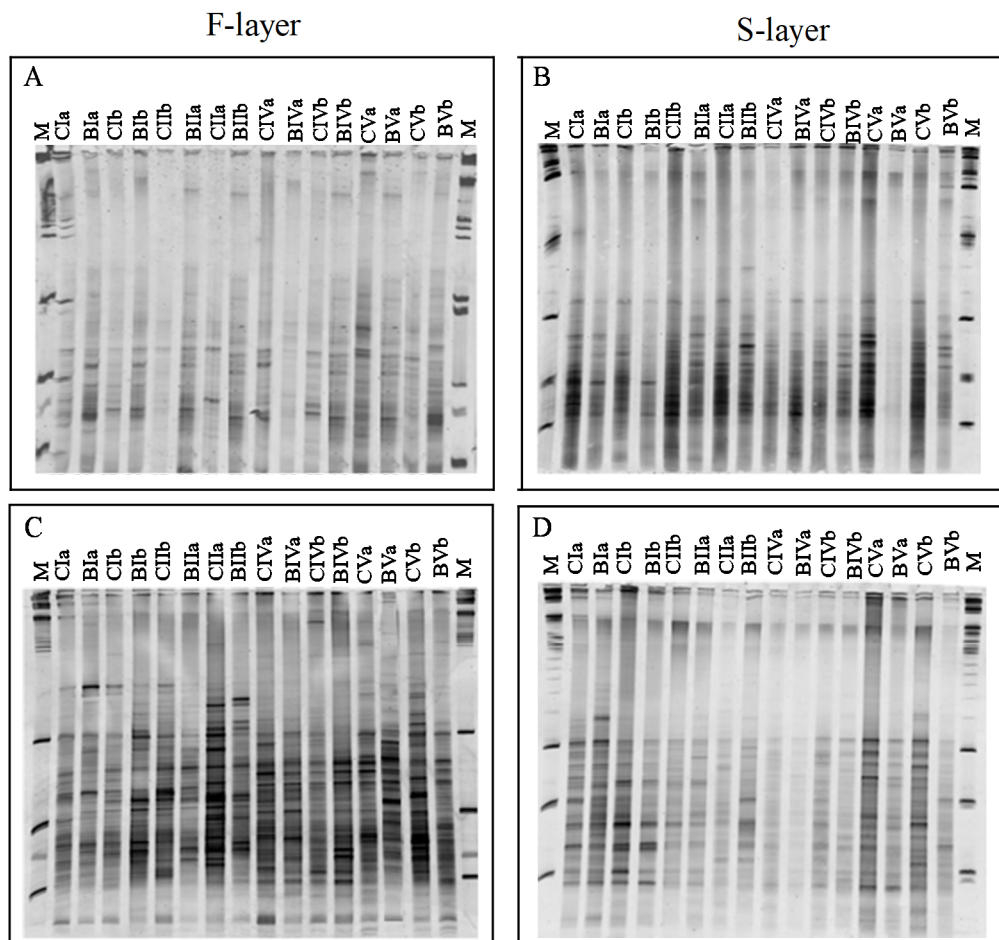


Figure 4. Genetic fingerprints of the bacterial community in *P. pinea* (A,B) and *P. pinaster* (C,D) plantations. For both plantation, two field replicates (a,b) for each experimental condition were analyzed at different times after prescribed fire (I, 3h; II, 33 d; IV, 189 d; and V, 363 d for *P. pinea* plantation and I, 3h; II, 32 d; IV, 182 d; and V, 371 d for *P. pinaster* plantation, respectively). M, marker Mass Ruler DNA Ladder mix.

Richness, Shannon, and evenness indexes (Table 4) were generally significantly ($p < 0.05$) higher in *P. pinaster* than in *P. pinea* plantation (except for soil richness), in line with C_{mic} and respiration trend ($p < 0.05$); in contrast, DNA amount was lower ($p < 0.05$) in the F-layer of *P. pinaster* plantation than in *P. pinea* plantation (Figures 2 and 3). *P. pinaster* plantation also showed significantly ($p < 0.01$) higher values of C_{org} and total N content, in the S-layer, and of C_{ext} and water content, in both F- and S-layer as compared with *P. pinea* plantation (Tables 2 and 3).

Two-way ANOVA analysis of the whole dataset showed that the sampling time was more critical than the burning treatment towards most variables considered (Tables 2 and 3, Figures 2 and 3), except for genetic ones that were generally unaffected by either of them (Table 4). The water content significantly changed with sampling time in both F- and S-layer (Table 3) and this in turn affected most other investigated variables. Indeed, extractable C (C_{ext}), microbial biomass, fungal mycelium of both F- and S-layer were generally positively correlated with the water content ($0.30 < r < 0.91$, $p < 0.05$, in *P. pinea* plantation, and $0.57 < r < 0.89$, $p < 0.001$, in *P. pinaster* plantation). Moreover, limited to the F-layer, total organic C (C_{org}) was positively correlated with water content in both plantations ($r = 0.32$, in *P. pinea*; $r = 0.26$, in *P. pinaster*, $p < 0.05$) and respiration was correlated with water content only

in *P. pinaster* plantation ($r = 0.33$, $p < 0.01$). In addition, soil mineral N ($\text{NH}_4^+\text{-N}$ and $\text{NO}_3^-\text{-N}$) was generally positively correlated with soil water content in both plantations ($0.27 < r < 0.49$, $p < 0.05$).

Table 4. Mean (\pm standard deviation) values of richness, Shannon, and evenness indexes in control and burned plots of *P. pinea* (A) and *P. pinaster* (B) plantations at different times after burning. Results of two-way ANOVA for burning and sampling time were reported at the bottom of A and B, respectively. N.S., nonsignificant.

(A) Time after Burning (Sampling Date)	F-Layer			S-Layer		
	Richness (Band Number)	Shannon Index	Evenness Index	Richness (Band Number)	Shannon Index	Evenness Index
3 h (12 March 2014)						
Control	27.0 (± 0.0)	3.1 (± 0.1)	0.96 (± 0.00)	39 (± 0.0)	3.3 (± 0.1)	0.91 (± 0.02)
Burned	27.5 (± 2.1)	3.1 (± 0.0)	0.92 (± 0.02)	36 (± 4.2)	3.3 (± 0.1)	0.93 (± 0.00)
33 d (14 April 2014)						
Control	29.0 (± 2.8)	3.2 (± 0.1)	0.94 (± 0.03)	39.0 (± 4.2)	3.4 (± 0.2)	0.93 (± 0.02)
Burned	26.5 (± 4.9)	3.0 (± 0.1)	0.94 (± 0.00)	35.0 (± 1.4)	3.4 (± 0.0)	0.94 (± 0.02)
189 d (17 September 2014)						
Control	24.0 (± 4.2)	3.0 (± 0.1)	0.96 (± 0.00)	39.0 (± 1.4)	3.5 (± 0.1)	0.95 (± 0.05)
Burned	25.5 (± 9.2)	3.0 (± 0.4)	0.93 (± 0.02)	33.0 (± 0.0)	3.3 (± 0.1)	0.94 (± 0.02)
363 d (10 March 2015)						
Control	21.5 (± 0.7)	2.9 (± 0.0)	0.94 (± 0.01)	39.5 (± 3.5)	3.4 (± 0.0)	0.93 (± 0.01)
Burned	22.0 (± 2.8)	2.7 (± 0.1)	0.90 (± 0.07)	38.0 (± 0.0)	3.4 (± 0.0)	0.94 (± 0.01)
Burning	N.S.	N.S.	N.S.	$P < 0.05$	N.S.	N.S.
Sampling time	N.S.	N.S.	N.S.	N.S.	N.S.	N.S.
(B) Time after Burning (Sampling Date)	F-layer			S-layer		
	Richness (band number)	Shannon index	Evenness index	Richness (band number)	Shannon index	Evenness Index
3 h (21 March 2014)						
Control	39.5 (± 2.1)	3.6 (± 0.1)	0.97 (± 0.02)	40.0 (± 5.7)	3.5 (± 0.1)	0.96 (± 0.00)
Burned	36.5 (± 0.7)	3.5 (± 0.1)	0.96 (± 0.01)	37.0 (± 1.4)	3.5 (± 0.1)	0.96 (± 0.02)
32 d (22 April 2014)						
Control	38.0 (± 4.2)	3.5 (± 0.1)	0.97 (± 0.01)	36.0 (± 2.8)	3.4 (± 0.2)	0.95 (± 0.03)
Burned	36.0 (± 2.8)	3.5 (± 0.1)	0.97 (± 0.01)	38.0 (± 1.4)	3.5 (± 0.0)	0.95 (± 0.01)
182 d (19 September 2014)						
Control	44.0 (± 2.82)	3.7 (± 0.1)	0.98 (± 0.01)	38.0 (± 7.1)	3.5 (± 0.1)	0.96 (± 0.01)
Burned	41.5 (± 3.5)	3.6 (± 0.1)	0.96 (± 0.01)	29.0 (± 7.1)	3.2 (± 0.2)	0.96 (± 0.01)
371 d (27 March 2015)						
Control	43.5 (± 0.7)	3.7 (± 0.0)	0.97 (± 0.00)	43.5 (± 2.1)	3.7 (± 0.0)	0.98 (± 0.00)
Burned	42.0 (± 0.0)	3.5 (± 0.1)	0.94 (± 0.01)	42.0 (± 5.7)	3.6 (± 0.1)	0.96 (± 0.00)
Burning	N.S.	N.S.	N.S.	N.S.	N.S.	N.S.
Sampling time	$p < 0.05$	N.S.	N.S.	N.S.	N.S.	N.S.

Bacterial diversity variables (richness, Shannon index, and evenness) generally did not change during the study period and were not correlated with microbial growth (DNA amount, microbial C, and fungal mycelium) and metabolism (respiration, $q\text{CO}_2$, and CMR) in *P. pinea* plantation. On the contrary, significant correlations among bacterial diversity and other microbial variables were observed in the *P. pinaster* plantation. In particular, richness and Shannon index were positively correlated with C_{mic} in the F-layer, and with respiration and CMR in the S-layer ($0.52 < r < 0.65$; $p < 0.05$). Surprisingly, richness in the F-layer was negatively correlated with respiration, $q\text{CO}_2$, CMR, and fungal fraction of C_{mic} ($-0.59 < r < -0.77$, $p < 0.05$).

4. Discussion

4.1. Effect of PB on Soil Microbial Community

As far as we are aware, this is the first study investigating the effects of prescribed burning (PB) on microbial biomass, activity and bacterial genetic diversity in pine plantations in a Mediterranean area. By comparing burned and control plots at each sampling time, we observed that the PB treatment in the *P. pinea* plantation did not affect the microbial community of either F- or S-layer (except for richness in soil). In contrast, a decrease in microbial biomass (C_{mic}), DNA amount, fungal mycelium, respiration, and C mineralization rate (CMR) was observed in the *P. pinaster* plantation generally until 32 d after treatment (in both layers or only one of the two). The more marked effect of prescribed burning in the *P. pinaster* plantation is consistent with higher maximum temperature and longer residence time above 100 °C during the treatment, most likely reflecting higher fire intensity due to more abundant litter fuel as compared with the *P. pinea* stand. Alcañiz et al. [30] reported that the effects of prescribed burning on soil biological properties depended on fire characteristics (intensity, residence time, and severity). Similarly, other authors observed that fire effect on soil microorganisms relied on temperatures at which they were exposed [35,65]. A decrease in fungal mycelium [13] and ATP content [66] with increasing fire severity has also been observed after experimental fires in an Italian Mediterranean maquis.

The effect of PB on soil microbial community in the *P. pinaster* plantation could be also due to indirect fire effect, such as the significant decrease of soil water content and C_{ext} concentrations in F- and S-layers of burned area, generally not observed in *P. pinea* plantation.

In this study, PB treatments were found to affect the soil microbial community to a lesser extent than that of sampling time. Indeed, two-way ANOVA showed that most variables depended on sampling time, only a few being also affected by fire treatment. Crucially, temporal variation due to rainfall seasonality largely overlapped the effect of PB in either plantation. As a matter of fact, soil biological activity in Mediterranean areas is primarily affected by yearly fluctuation in water availability [67]. In line with this, most microbial variables considered in this study were positively correlated to the water content.

Our results are consistent with former research showing that repeated PB in a *P. halepensis* plantation in Southern Italy had only short-term or no effects on chemical and microbial properties of the fermentation layer and the 5 cm soil layer underneath, with a superimposed predominance of the water factor [31]. Likewise, no or minimal effects of PB on soil physical and chemical properties were observed in a short-grass plain in Texas [37], a *P. pinaster* plantation in Portugal [68] and a *P. palustris* forest in South Carolina [69].

Total organic C did not change significantly in burned plots as compared with controls, while F-layer extractable C (limited to 89 and 363 days) and soil total N and mineral N significantly increased in burned plots of *P. pinea* plantation, where fire intensity was relatively low. This possibly reflects incorporation of nitrogen-enriched ashes produced by combustion at low temperature [30]. An increase in ammoniacal N has been reported in a Texas grassland after PB [37]. Benefits from higher C and N contents could balance negative effects of fire, thus, accounting for little or no damage to the microbial community. Indeed, in *P. pinea* plantation, soil DNA and fungal mycelium were positively correlated with NH_4^+ -N concentration; moreover, soil respiration, qCO_2 , and CMR were positively correlated with NO_3^- -N concentration.

4.2. Interrelationships among Variables

Bacterial genetic diversity was evaluated as richness, Shannon index, and evenness of bacteria after DNA extraction and amplification of a specific segment of bacterial 16S rDNA [60]. The values of DNA yield for the F-layer of *P. pinea* (44 to 79 mg kg⁻¹) and *P. pinaster* (25 to 58 mg kg⁻¹) plantations and for the S-layer (respectively, 13 to 51 mg kg⁻¹ in *P. pinea*, and 11 to 38 mg kg⁻¹, in *P. pinaster*) mostly fell within the range (0.1 to 41.8 mg kg⁻¹) reported in 2150 French soils [52], only occasionally being slightly above. The richness values found in our study systems were comparable with those

reported in soils from *P. nigra* and *Abies alba* Italian forests [70] and in a *P. pinaster* Spanish forest with an understory of Mediterranean bushes [71].

Whereas wildfire can cause changes in the soil bacterial genetic diversity, as demonstrated in a Spanish *P. pinaster* forest [71], our study showed that generally PB did not affect bacterial genetic diversity in either plantation, except for richness in soil of *P. pinea* plantation. Because bacterial richness in the above-lying F-layer was not affected by treatment, the observed effect on soil suggests the involvement of other factors (not investigated here) that co-varied with the burning factor. Other authors found no or temporary variations in microbial community structure after prescribed burning [37,72]. On the contrary, experimental summer fire in a Mediterranean maquis caused decreases in functional diversity [56], fungal species density [73], and fungal fraction of microbial C [13], although an increase in culturable total, xerotolerant, and heat-stimulated fungi was also observed [74]. In this study, an increase in the fungal fraction of microbial C was only found in the S-layer in *P. pinaster* plantation and was restricted to the first month after treatment. In combination with a parallel decrease in C_{mic} and fungal mycelium, this suggests that fire negatively affected bacteria more than fungi in the soil of the *P. pinaster* stand.

Unchanged genetic bacterial diversity in burned plots in *P. pinea* plantation is an expected result, because PB had relatively low intensity and no other microbial variable changed appreciably after exposition to fire in this system. This is consistent with the lack of effects in this plantation also on tree growth [24] and plant species richness (Esposito et al., unpublished data). The lack of effect on bacterial diversity in the *P. pinaster* plantation, instead, is more surprising because a higher fire intensity in this system caused significant changes in other microbial variables, generally persisting for a month after fire, and reduced microbial efficiency in C use as shown by higher values of soil qCO_2 . A significant increase in soil qCO_2 after wildfire and experimental summer fire has also been reported in Mediterranean maquis [13,75]. These observations fit well with the general tendency of microorganisms to increase CO_2 -producing catabolic activity and reduce growth-sustaining anabolism when exposed to stressing conditions [6,50,76].

The lack of PB effects on soil bacterial diversity and the temporary effects on soil microbial biomass and activity in *P. pinaster* plantation demonstrated that also higher intensity PB had a low impact on soil. On the other hand, in this plantation PB did not affect negatively the tree growth [42] nor the plant species richness. Indeed, 20 months after PB treatment plant species richness did not change between control and burned plots (6.4 ± 1.9 vs. 6.9 ± 1.3 , as measured in 78.5 m^2 plots, Stinca et al., unpublished results).

No correlation was observed in *P. pinea* plantation among bacterial genetic diversity and microbial biomass and activity, whereas, in *P. pinaster* plantation, richness, and Shannon index were positively correlated with C_{mic} in the F-layer, and with respiration and CMR in the S-layer. On the other hand, richness was negatively correlated with microbial activity and fungal fraction of C_{mic} in the F-layer. These contradictory relationships suggest the participation of factors other than those considered in the present study. Nannipieri et al. [8] suggested that the lack of correlation between microbial diversity and a fundamental index of microbial activity such as organic matter decomposition could be explained by functional redundancy of the microbial community. As a matter of fact, the reduction and disappearance of certain microbial entities could have little effect on soil functions if others persist that can perform the same functions.

Relatively to intra-ecosystem variability, a greater variability in bacterial genetic diversity was observed when comparing the two pine plantations (about 40 km apart) to each other, demonstrating that these indices are sensitive to variation of environmental factors (such as those site dependent). Indeed, as compared with *P. pinea* plantation, the *P. pinaster* plantation generally showed significantly higher values of richness, Shannon index, and evenness in both the F- and S-layer, which fits well with higher values of C_{mic} and respiration. Surprisingly, an opposite trend was observed for soil DNA amount. In *P. pinaster* plantation, soil microorganisms could benefit from significantly higher soil organic C and total N content as well as higher C_{ext} and water content in both F- and S-layer.

This confirms the importance of water as a regulating factor for soil microbial community in the Mediterranean environment [66]. As compared with *P. pinea* plantation, *P. pinaster* plantation had a lower undergrowth plant species richness, as well as lower shrub and herbs cover and tree high, suggesting that microbial community was affected by soil chemical properties more than plant species richness and vegetation structure.

5. Conclusions

The application of PB for fire hazard reduction in two pine plantations located in protected areas of Southern Italy exposed to high fire risk achieved the objective of reducing the litter fuel and interrupting its continuity without causing persistent negative effects on the soil, as well as on tree growth and vegetation. In this study a temporary negative effect on microbial biomass and activity was observed in the *P. pinaster* plantation only, where higher fire intensity occurred as compared with *P. pinea* plantation. However, in both plantations the magnitude of fire-related changes was by far smaller than the spectrum of changes associated with the sampling time, confirming our first hypothesis.

Moreover, our results confirmed our second hypothesis that microbial diversity is a powerful tool that effectively integrates microbial growth and activity data in the study of fire effects on soil microbial communities. Indeed, the absence of changes in bacterial genetic diversity in burned plots suggested that fire caused no important damage to the structure of microbial community living immediately below the litter layer even after relatively intense burning in *P. pinaster* plantation, thus explaining the fast recovery of microbial growth and activity in the aftermath.

Our results confirm that the application of prescribed burning in the studied plantations under the prescriptions adopted (see Table 1) could be a safe practice reducing forest fire hazard without causing relevant damages.

Questions open to future research, include: (i) to verify if the response to PB observed in pine plantations can be generalized to other Mediterranean ecosystems, (ii) to test fire intensity and severity thresholds above which marked changes in soil microbial activity occur, and (iii) to know if specific bacterial or fungal species could be damaged or stimulated by PB treatment.

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