



Invasion of Native Riparian Forests by *Acacia* Species Affects In-Stream Litter Decomposition and Associated Microbial Decomposers

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Abstract

The invasion of native riparian forests by exotic tree species can lead to profound changes in the ecological integrity of freshwater ecosystems. We assessed litter decomposition of native (*Alnus glutinosa* and *Quercus robur*) and invasive (*Acacia melanoxylon* and *Acacia dealbata*) tree species, and associated microbial activity and community structure, after being immersed for conditioning in 3 reference and 3 “invaded” streams in Serra da Lousã (central Portugal) and used in microcosms simulating stream conditions. Litter decomposition differed among species, in the order: *Al. glutinosa* > *Q. robur* > (*Ac. melanoxylon* ~ *Ac. dealbata*). *Alnus glutinosa* litter decomposed faster probably because it was soft and had high nitrogen concentration for decomposers. *Quercus robur* litter decomposed slower most likely because it was tough and had high polyphenol and low nitrogen concentrations. *Acacia melanoxylon* litter was the toughest and had a thick cuticle that likely acted as a physical barrier for microbial colonization. In *Ac. dealbata*, the small-sized leaflets and high lignin concentration may have limited microbial litter decomposition. Litter decomposition was faster in “invaded” streams, probably because they were N-limited and increases in nitrogen concentration in water, promoted by *Acacia* species invasion, stimulated microbial activity on litter. The aquatic hyphomycete community structure differed among litter species and between stream types, further suggesting that microbes were sensitive to litter characteristics and water nutrient concentrations. Overall, the invasion of native riparian forests by *Acacia* species may affect microbial decomposer activity, thus altering important stream ecosystem processes, such as litter decomposition and nutrient cycles.

Keywords *Acacia dealbata* · *Acacia melanoxylon* · Aquatic hyphomycetes · Forest invasion · Leaf decomposition · Nitrogen-fixing species

Introduction

The invasion of native forests by exotic plant species is occurring worldwide and is expected to increase due to human activities (e.g., agriculture, urbanization) and climate change (e.g., increased temperature, altered precipitation) [1, 2]. Plant

invasions are expected to change native forest composition and alter the diversity of plant functional traits [2]. Therefore, plant invasions can have profound effects on detrital ecosystems, which depend on plant matter inputs [2–4].

In forest streams, the plant matter supplied by the riparian vegetation (e.g., leaves, wood) represents a major source of organic carbon for aquatic food webs [5]. In these ecosystems, plant litter decomposition is a fundamental ecosystem process mediated by the activities of microbes, namely fungi and bacteria, and invertebrate detritivores [6]. Fungi, particularly aquatic hyphomycetes, and bacteria produce a large variety of extracellular enzymes that promote leaf maceration and increase leaf palatability to invertebrate detritivores [6, 7]. Aquatic hyphomycetes are key decomposers of litter particularly in the initial stages of the process, while bacteria only increase their contribution when litter has been partially

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broken down [8]. Altogether, aquatic decomposers mineralize the organic carbon from litter, convert it into biomass, and promote the release of dissolved and fine organic particles that cause leaf mass loss [7]. Microbial decomposers respond to changes in litter characteristics and nutrient concentrations in stream water. Microbes generally colonize litter of high nutritional quality (i.e., soft litter, with high nutrient concentration and low concentrations of structural and secondary compounds) and decompose it faster than more recalcitrant litter [9–12], and have higher activity under nutrient-rich conditions [10, 13, 14]. Additionally, aquatic hyphomycete community structure is sensitive to resource diversity (e.g., leaf litter), being more diverse when resource diversity (i.e., riparian vegetation and litter inputs) is higher [15–17]. Also, some aquatic hyphomycete species can occur as endophytes, and thus, increases in riparian diversity may promote an increase in aquatic hyphomycete diversity due to host specificity between endophytes and trees [18].

Over the past 250 years, many Australian *Acacia* species have been introduced into several regions of the world [19], mainly for wood and timber production, to stabilize dunes and to prevent soil erosion [20]. Specific traits such as resistant seeds that quickly germinate, capacity to fix atmospheric nitrogen (N), high capacity to adapt to different environmental conditions, rapid growth rate, and high accumulation of biomass are responsible for their successful invasive behavior [20, 21]. In Portugal, *Acacia* species cover at least ~ 17,000 ha [22], and it is forbidden to further introduce and use the entire *Acacia* genus for afforestation [23]. *Acacia* species are among the 10 most problematic invasive exotic species in Portugal and among the 30 most problematic in Spain and France [24]. Invasion of native forests by *Acacia* species is expected to affect the characteristics of litter inputs to streams as diverse deciduous forests with low abundances of N-fixers are being replaced by monospecific stands of evergreen N-fixing species [20]. Moreover, N concentrations in stream water were expected to increase as found for streams with riparian areas invaded by other exotic N-fixing species [25–29]. Given the sensitivity of microbial decomposers to litter characteristics and dissolved nutrient concentrations, changes in these parameters can have strong effects on litter decomposition in “invaded” streams.

Currently, in central Portugal, many native riparian deciduous forests dominated by *Castanea sativa* Mill. and *Quercus robur* L. are being invaded by exotic N-fixing species (mostly *Acacia dealbata* Link. and to a lesser extent *Acacia melanoxylon* R. Br.). Here, we examine the effects of litter identity and quality (native species: *Alnus glutinosa* (L.) Gaertn and *Q. robur*; invasive species: *Ac. melanoxylon* and *Ac. dealbata*) and dissolved nutrient concentrations (nutrient levels from reference and “invaded” streams) on litter decomposition and associated microbial decomposers in laboratory microcosms simulating stream conditions. We predict that (i)

litter with high nutritional quality (soft, high initial N concentration) will have higher microbial activity and faster litter decomposition; (ii) microbial activity will be higher, and litter decomposition will be faster in “invaded” than reference streams due to a stimulation of microbial activity by increased N concentrations in stream water; and (iii) litter in reference streams will have higher aquatic hyphomycete species richness due to a higher inoculum potential as a result of higher resource diversity.

Material and Methods

Study Region

Six small forest streams (< 4 m wide), located in Serra da Lousã (central Portugal), were selected for this experiment (Table 1). Reference streams (Cerdeira, Candal, and Maior) run through native forests mainly composed of *C. sativa* and *Q. robur* trees, while “invaded” streams (Sotão, Fiscal, and Piedade) run through forests heavily invaded by *Acacia* species (94–100% cover in the riparian area defined as 50 m to each side of the stream and 250 m long is composed mostly by *Ac. dealbata* (> 90%) and to a lesser extent by *Ac. melanoxylon* (< 10%). All streams run on schist bedrock. Stream basins have low human occupation, and agricultural activity is nearly absent (< 0.5% of basin area is occupied by agriculture).

Litter Species and Initial Chemical Characteristics

Litter of two native species, *Al. glutinosa* and *Q. robur*, and of two invasive species, *Ac. melanoxylon* and *Ac. dealbata*, commonly found in the riparian vegetation of streams flowing through native and “invaded” forests of central Portugal, respectively, were selected for this study. The native species were selected among the dominant riparian species to represent extremes in terms of leaf litter N concentration [30]. *Alnus glutinosa* is an actinorrhizal species that establishes symbiosis with the N-fixing actinobacteria *Frankia alni*, thus acting as an N-fixing tree species, while *Q. robur* is not an N-fixer. *Acacia* species are N-fixers, but while *Ac. melanoxylon* has phyllodes (i.e., leaf-like structures derived from modified petioles), *Ac. dealbata* has bipinnate leaves with a large number of small leaflets (2–5 mm long and 0.4–0.7 mm wide).

Litter was collected in Serra da Lousã, central Portugal, in autumn 2016. Litter of *Al. glutinosa*, *Q. robur*, and *Ac. melanoxylon* were collected just after abscission from the ground in small tree stands dominated by the target species, where trees were exposed to similar environmental and edaphic conditions and appeared to have similar age and vigor. *Acacia dealbata* litter was collected directly from a few trees of similar age and vigor in an *Ac. dealbata* dominated stand.

Table 1 Location and physical and chemical water characteristics of the six study streams in Serra da Lousã (central Portugal)

Stream type	Latitude (N)	Longitude (W)	Elevation (m)	Temperature (°C)	pH	Dissolved O ₂ (mg L ⁻¹)	Conductivity (μS cm ⁻¹)	NO ₂ ⁻ -N (μg L ⁻¹)	NO ₃ ⁻ -N (μg L ⁻¹)	NH ₄ ⁺ -N (μg L ⁻¹)
Reference										
Cerdeira	40° 05' 23.1"	8° 12' 05.0"	529	11.5 ± 1.08	7.3 ± 0.09	10.8 ± 0.40	35.3 ± 2.78	0.5 ± 0.03	3.7 ± 0.23	20.1 ± 4.00
Maior	40° 07' 53.3"	8° 11' 40.7"	195	13.1 ± 1.35	7.4 ± 0.09	11.3 ± 0.09	43.3 ± 4.85	0.5 ± 0.03	4.3 ± 2.17	20.3 ± 4.09
Candal	40° 04' 54.1"	8° 12' 16.6"	634	11.0 ± 0.85	7.3 ± 0.07	10.6 ± 0.20	28.5 ± 0.87	0.5 ± 0.10	3.3 ± 0.50	22.9 ± 1.74
“Invaded”										
Sotão	40° 07' 54.1"	8° 09' 08.3"	373	12.2 ± 0.79	7.4 ± 0.07	10.8 ± 0.32	59.0 ± 1.58	0.6 ± 0.09	29.6 ± 6.10	14.9 ± 3.00
Fiscal	40° 06' 40.2"	8° 13' 35.1"	329	12.0 ± 0.80	7.3 ± 0.02	10.9 ± 0.36	65.5 ± 2.72	0.5 ± 0.06	13.0 ± 0.81	21.6 ± 1.57
Piedade	40° 05' 52.6"	8° 14' 11.5"	250	10.0 ± 1.09	7.3 ± 0.14	10.7 ± 0.32	62.3 ± 1.80	0.6 ± 0.08	13.2 ± 1.08	17.7 ± 0.02
Average										
Reference				11.8 ± 0.64 ^a	7.3 ± 0.04 ^a	10.8 ± 0.19 ^a	35.7 ± 2.49 ^a	0.5 ± 0.03 ^a	3.8 ± 0.61 ^a	21.1 ± 1.64 ^a
“Invaded”				11.4 ± 0.56 ^a	7.3 ± 0.05 ^a	10.8 ± 0.18 ^a	62.3 ± 1.35 ^b	0.6 ± 0.05 ^a	18.6 ± 3.83 ^b	18.1 ± 1.50 ^a

Values are means ± SE ($n = 2-4$). Stream types with different letters are significantly different (one-way ANOVA followed by Tukey HSD test, $p < 0.05$)

Litter of each species was pooled together, air-dried at room temperature, and stored in the dark until used.

Three sets of air-dried litter from each species (randomly selected from each species litter pool) were oven-dried (105 °C, ± 48 h) and ground to a fine powder (< 0.5 mm size; Retsch, ZM 100 Ultra Centrifugal Mill, Haan, Germany). Then, the powder was oven-dried (105 °C, ± 48 h) and analyzed to determine the initial polyphenols [31], lignin [32], phosphorus [33], carbon (C), and nitrogen (N) (Thermo Fisher Scientific Inc., CNH auto analyzer IRMS Thermo Delta V advantage with a Flash EA, 1112 series, Waltham, USA) concentrations of each species prior to the beginning of the experiment. Results were expressed as percentage of dry mass (DM).

Microbial Colonization of Litter

Air-dried litter of *Al. glutinosa*, *Q. robur*, *Ac. melanoxyton*, and *Ac. dealbata* were enclosed individually in fine-mesh bags (0.5 mm mesh size; 5 litter bags per litter species) ad libitum, and litter bags were immersed in the three reference and the three “invaded” streams on March 2017 to allow microbial colonization. On the same occasion, conductivity, water temperature, pH, and dissolved O₂ concentration were measured in situ with field probes (WTW, Weilheim, Germany). Stream water (± 8 L) was also collected in acid-washed plastic bottles from each stream, transported in a cooler to the laboratory, filtered through glass microfiber filters (47 mm diameter, 0.7 μm pore size; Whatman GF/F, GE Healthcare UK Limited, Little Chalfont, UK), and frozen at -20 °C until needed. Stream water was used for the microcosm experiment and for determination of nutrient

concentrations. Nitrite, nitrate, and ammonium concentrations were determined by the colorimetric method (AA3 Bran + Luebbe autoanalyzer; SEAL Analytical, Norderstedt, Germany).

Microcosm Setup

After 1 week of colonization in the streams, litter bags were collected, placed individually in zip-lock bags, and transported in a cooler to the laboratory. In the laboratory, litter was rinsed with distilled water. Discs from *Al. glutinosa*, *Q. robur*, and *Ac. melanoxyton* litter were cut with a cork borer (12-mm diameter; avoiding the main vein for *Al. glutinosa* and *Q. robur*), and leaflets from *Ac. dealbata* were detached from the secondary veins. Litter discs (20 per microcosm) and portions of leaflets were wet-weighed (± 0.1 mg) and transferred to 100-mL Erlenmeyer flasks containing 40 mL of filtered water from the stream of origin. Microcosms were stored on an orbital shaker (100 rotations minute⁻¹; GFL 3017, Burgwedel, Germany) at 18 °C, under a 12-h light and 12-h dark regime. Each stream within each stream type acted as a replicate, i.e., Cerdeira, Maior, and Candal were the three replicate streams for the reference stream type and Sotão, Fiscal, and Piedade were the three replicate streams for the “invaded” stream type (4 litter species × 2 stream types × 3 replicate streams × 4 sampling dates = 96 microcosms) (Fig. 1). The water in the microcosms was renewed twice a week using filtered water from the stream of origin. After 14, 31, 49, and 70 days of incubation in the laboratory, three microcosms per treatment (i.e., one flask per litter species per stream) were sacrificed for the determination of microbial respiration, fungal biomass, conidia

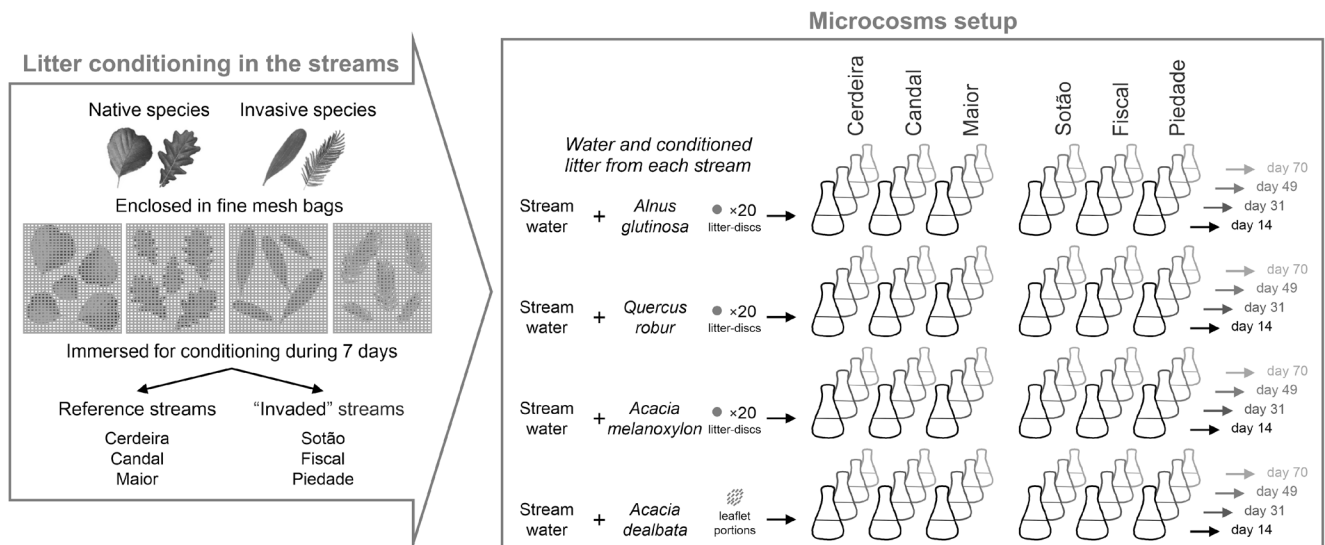


Fig. 1 Experimental design showing that litter of *Alnus glutinosa*, *Quercus robur*, *Acacia melanoxylon*, and *Acacia dealbata* was enclosed in fine-mesh bags and immersed for conditioning during 7 days in three reference (Cerqueira, Candal and Maior) and three “invaded” streams (Sotão, Fiscal, and Piedade), in Serra da Lousã (central Portugal). Sets of 20 discs were cut from *Al. glutinosa*, *Q. robur*, and

Ac. melanoxylon litter and portions of leaflets were detached from *Ac. dealbata* litter and incubated in flasks containing water from the stream of origin. Three microcosms per treatment were sacrificed after 14, 31, 49, and 70 days of incubation for mass remaining and microbial determinations

production by aquatic hyphomycetes, litter toughness, and litter mass loss (see below).

Microbial Respiration

On each sampling date, a subset of five litter discs or a portion of leaflets from each sacrificed microcosm was used to determine the microbial oxygen consumption rates using a closed six-channel dissolved oxygen measuring system (Strathkelvin 929 System, North Lanarkshire, UK). A saturated solution of sodium sulfite in 0.01 M sodium borate (0% O_2 ; prepared immediately before use) and an O_2 -saturated water sample from the stream of origin (100% O_2 , kept at 18 °C) were used to calibrate the oxygen electrodes. Then, litter discs and portions of leaflets were incubated in 3-mL chambers with the corresponding saturated stream water kept at 18 °C. Chambers without plant material were used as controls. The system was left running for ± 45 min. Then, litter discs and leaflets were removed from the chambers and used to determine litter toughness (see below) and litter discs DM. Oxygen consumption rates were determined by the difference in the O_2 concentration in the sample and the control, at the beginning and the end of a 15-min interval during which O_2 consumption over time was linear (taken between 10 and 40 min of incubation). Results were expressed as $mg\ O_2\ g^{-1}\ DM\ h^{-1}$.

Fungal Biomass

Another subset of five litter discs and a portion of leaflets were frozen at $-20\ ^\circ C$, lyophilized overnight, weighed (± 0.1 mg)

to determine DM and used for ergosterol extraction [31, 34]. Litter discs and leaflets were placed in tightly closed tubes with 10 mL of KOH in methanol ($8\ g\ L^{-1}$). Lipids were extracted in an 80 °C water bath for 30 min, purified by solid phase extraction (Waters Sep-Pak Vac RC tC18 cartridges; Waters Corp., Milford, USA) and eluted with isopropanol. Ergosterol was quantified by high-performance liquid chromatography (HPLC; Dionex DX-120, Sunnyvale, CA, USA) using a Thermo Scientific Synchronis C18 column (250×4 mm, 5 μm particle size; Thermo, Waltham, MA, U.S.A.) and HPLC-grade methanol as the mobile phase ($1.4\ mL\ min^{-1}$ at 33 °C). Ergosterol was detected at 282 nm, quantified based on a standard curve of ergosterol in isopropanol, and converted into mycelial biomass assuming $5.5\ \mu g\ ergosterol\ mg^{-1}\ fungal\ DM$ [34]. Results were expressed as $mg\ fungal\ DM\ g^{-1}\ litter\ DM$.

Conidia Production by Aquatic Hyphomycetes

On each sampling date, the conidial suspension from each sacrificed microcosm was saved into 50-mL Falcon tubes, fixed with 2 mL of 37% formalin, and stored in the dark until processed. Suspensions were then mixed with 100 μL of 0.5% Triton X-100, and appropriate volumes of each sample were filtered through cellulose nitrate filters (25 mm diameter, 5 μm pore size; Sartorius Stedim Biotech GmbH, Goettingen, Germany). Filters were stained with 0.05% cotton blue in 60% lactic acid, and conidia were counted and identified under a microscope (200 \times magnification; Leica, DM1000, Wetzlar, Germany) [31]. Sporulation rates were expressed as

the number of conidia released mg^{-1} litter DM day^{-1} and species richness as the number of species sample^{-1} .

Litter Toughness

Litter toughness was determined for *Al. glutinosa*, *Q. robur*, and *Ac. melanoxyton* litter discs using a penetrometer [31]. Toughness was measured as the mass (± 0.01 g) required for a pin, from a 1.55-mm diameter iron rod, to penetrate through the litter, avoiding major veins [31]. Toughness was not determined for *Ac. dealbata* litter since it was not possible to apply this method to the small leaflets. Litter toughness was used as a surrogate for enzymatic maceration of litter and expressed as percentage of initial litter toughness. Initial litter toughness was determined applying the same method to litter discs extracted after the 1-week colonization period.

Litter Decomposition

The remaining 10 litter discs and portions of leaflets from each sacrificed microcosm were oven-dried at 105 °C for 24 h and weighed (± 0.1 mg). This DM was added to that of discs and portions of leaflets used for the determination of microbial respiration and fungal biomass to determine DM remaining. Litter DM remaining comprised the biomass of associated fungi. Results were expressed as percentage of initial DM. Initial DM was estimated by multiplying the initial wet mass by a conversion factor derived from an additional group of wet litter discs and leaflets from each stream. These samples were oven-dried at 105 °C for 24 h, weighed (± 0.1 mg) to determine the litter discs and leaflets initial DM, and the conversion factor was calculated as the ratio between the initial DM and the initial wet mass. Litter was incubated in the field for a short period, and it was carefully rinsed with distilled water before discs were cut or leaflets detached, and therefore, adherent sediments were not a concern and there was no need to estimate litter ash fraction.

Data Analysis

Data normality was checked by D'Agostino-Pearson or Kolmogorov-Smirnov test and homoscedasticity was checked by Bartlett or Levene's test. When data failed to meet one of these assumptions, they were transformed (e.g., fungal biomass) before analysis.

Water characteristics were compared between stream types, and litter initial chemical characteristics were compared among species by one-way analysis of variance (ANOVA), followed by Tukey's Honest Significant Difference (HSD) test when needed.

Decomposition rates (k , d^{-1}) were calculated assuming an exponential decay model, by linear regression of the \ln -transformed fraction of DM remaining against time (days), and a linear decay model, by linear regression of the fraction

of DM remaining against time (days). Since none of these models properly fits the data in all treatments, the effects of litter species and stream type on litter decomposition, given by the DM remaining at the last sampling date, were tested by two-way ANOVA, followed by Tukey's HSD test.

Fungal biomass (Box-Cox transformed to meet the assumptions of ANOVA), aquatic hyphomycete species richness and litter toughness were compared among litter species, stream type, and time (days) by three-way ANOVA, followed by Tukey's test. Microbial respiration rates and fungal sporulation rates were compared among litter species, stream type, and time (days) by three-way univariate permutational analysis of variance (PERMANOVA), since data failed to meet the ANOVA assumptions of homoscedasticity even after transformation. Univariate PERMANOVAs were run on Euclidean distance [35], and residuals were permuted under a reduced model of 9999 permutations, followed by pair-wise comparison tests [35]. Univariate PERMANOVA is equivalent to a standard ANOVA and can be used when data fail to meet the assumptions of ANOVA [35]. For each of the measured parameters, the square of the correlation ratio (η^2), i.e., the percentage of the variance accounted for each of the main effects and interactions, was calculated as $\text{SS}_{\text{effect}}/\text{SS}_{\text{total}} \times 100$ [36].

The structure of aquatic hyphomycete communities was assessed by cluster analysis, based on the Bray-Curtis similarity matrix of conidial production by aquatic hyphomycetes ($\log(x + 1)$ transformed). Aquatic hyphomycete communities were compared among litter species and between stream types by two-way multivariate PERMANOVA, followed by pair-wise comparison tests.

Cluster analysis, PERMANOVA and pair-wise tests were performed on PRIMER 6 (v6.1.16) and PERMANOVA+ (v1.0.6; Primer-E Ltd., Plymouth, UK). All other statistical analyses were performed on the STATISTICA 8 software (StatSoft Inc., Tulsa, Oklahoma, USA).

Results

Stream Water

During leaf colonization, stream water was cool, circumneutral, well-oxygenated, and had low conductivity and low nutrient concentrations (Table 1). Water characteristics were similar between stream types, except for conductivity and NO_3^- -N, which were significantly higher in "invaded" than in reference streams (one-way ANOVA, $p < 0.001$ and $p = 0.003$, respectively) (Table 1).

Litter Chemistry

Litter species differed in initial chemical characteristics (one-way ANOVA, $p < 0.05$), except for carbon concentration

(Table 2). *Quercus robur* litter had the highest polyphenols and phosphorus concentrations and C:N ratio, and the lowest N concentration (Table 2). *Acacia dealbata* litter had the highest N concentration, higher lignin concentration than *Q. robur*, and the lowest polyphenol concentration and C:N ratio together with *Al. glutinosa*. *Acacia melanoxylon* litter had the lowest phosphorus concentration (Table 2).

Litter Decomposition

Litter mass decreased during the incubation period for all species (Fig. 2a–d). After 70 days, litter mass remaining varied between 52 and 50% for *Al. glutinosa*, 69 and 64% for *Q. robur*, 79 and 77% for *Ac. melanoxylon*, and 82 and 77% for *Ac. dealbata*, in reference and “invaded” streams, respectively. Litter mass remaining differed significantly among litter species and between stream types (two-way ANOVA, $p < 0.001$ and $p = 0.021$, respectively), with no significant interaction between factors ($p = 0.631$) (Table S1). Litter mass remaining was significantly higher in reference than in “invaded” streams (two-way ANOVA, $p = 0.021$) and was in the order: *Al. glutinosa* < *Q. robur* < (*Ac. melanoxylon* ~ *Ac. dealbata*) (Tukey’s test, $p < 0.001$). Litter species accounted for a disproportionally higher variance in litter mass remaining ($\eta^2 = 92\%$) when compared with stream type ($\eta^2 = 2\%$) (Table S1).

Litter Toughness

Litter toughness remaining decreased mostly during the first 14 days of incubation for all species and after 70 days litter toughness varied between 25 and 21% for *Al. glutinosa*, 48 and 42% for *Q. robur*, and 55 and 60% for *Ac. melanoxylon*, in reference and “invaded” streams, respectively (Fig. 2e–g). Litter toughness differed significantly among litter species (three-way ANOVA, $p < 0.001$) (Table S2), with lower toughness for *Al. glutinosa*, followed by *Q. robur* and *Ac. melanoxylon* (Tukey’s test, $p < 0.001$). Litter species accounted for 59% of the variance in litter toughness (Table S2). Litter toughness did neither significantly differ

between stream types nor was there a significant interaction between litter species and stream type (three-way ANOVA, $p = 0.380$ and $p = 0.222$, respectively) (Table S2).

Microbial Respiration Rate

After 70 days of incubation, microbial respiration rates varied between 0.40 and 0.39 mg O₂ g⁻¹ DM h⁻¹ for *Al. glutinosa* litter, 0.16 and 0.26 mg O₂ g⁻¹ DM h⁻¹ for *Q. robur*, 0.12 and 0.15 mg O₂ g⁻¹ DM h⁻¹ for *Ac. melanoxylon*, and 0.48 and 0.55 mg O₂ g⁻¹ DM h⁻¹ for *Ac. dealbata*, in reference and “invaded” streams, respectively (Fig. 2h–k). Microbial respiration rates differed significantly among litter species (three-way univariate PERMANOVA, $p < 0.001$) and were significantly affected by the interaction between litter species and stream type ($p = 0.004$) (Table S3). Microbial respiration was higher in *Al. glutinosa* litter, followed by *Ac. dealbata* and *Q. robur*, and lower in *Ac. melanoxylon* (pair-wise test, $p < 0.002$). *Quercus robur*, *Ac. melanoxylon*, and *Ac. dealbata* litter supported higher microbial respiration in “invaded” than in reference streams (pair-wise test, $p = 0.013$, $p < 0.001$, and $p = 0.001$, respectively). Litter species accounted for a higher variance in microbial respiration ($\eta^2 = 46\%$) than the interaction between litter species and stream type ($\eta^2 = 4\%$) (Table S3).

Fungal Biomass

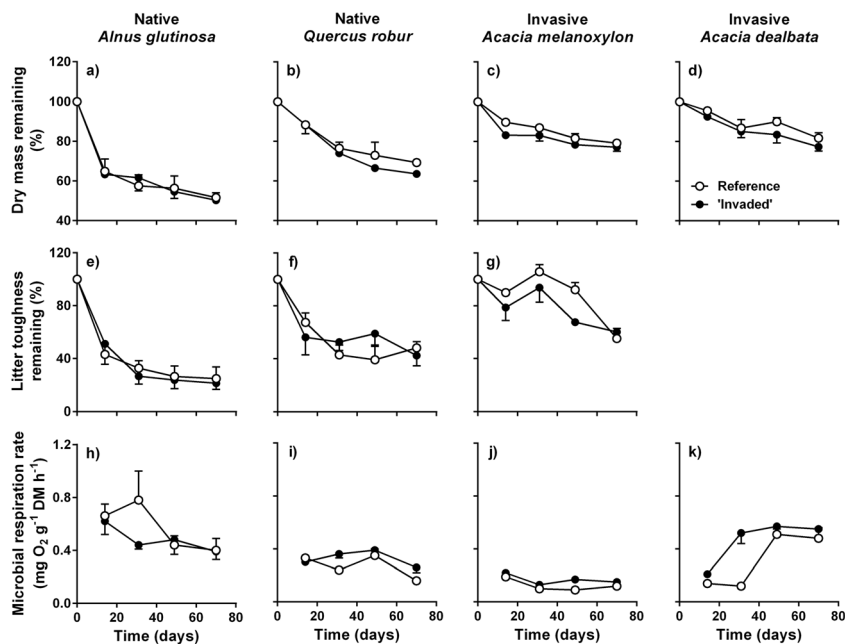
After 70 days of incubation, fungal biomass varied between 32 and 30 mg g⁻¹ DM for *Al. glutinosa* litter, 57 and 51 mg g⁻¹ DM for *Q. robur*, 23 and 24 mg g⁻¹ DM for *Ac. melanoxylon*, and 42 and 33 mg g⁻¹ DM for *Ac. dealbata*, in reference and “invaded” streams, respectively (Fig. 3a–d). Fungal biomass differed significantly among litter species (three-way ANOVA, $p < 0.001$) and was significantly affected by the interaction between litter species and stream type ($p < 0.001$) (Table S2). *Quercus robur* and *Ac. dealbata* litter had higher fungal biomass than *Al. glutinosa* and *Ac. melanoxylon* (Tukey’s test, $p < 0.001$). *Alnus glutinosa* and *Ac. dealbata* litter had higher fungal biomass in reference

Table 2 Initial chemical characteristics of native (*Alnus glutinosa* and *Quercus robur*) and invasive (*Acacia melanoxylon* and *Acacia dealbata*) species used in the microcosm experiment

Litter variables	<i>Alnus glutinosa</i>	<i>Quercus robur</i>	<i>Acacia melanoxylon</i>	<i>Acacia dealbata</i>
Polyphenols (%)	3.8 ± 0.7 ^{bc}	13.2 ± 0.1 ^a	4.8 ± 0.4 ^b	2.2 ± 0.1 ^c
Lignin (%)	38.5 ± 0.7 ^{ab}	34.1 ± 2.1 ^b	38.1 ± 1.2 ^{ab}	42.9 ± 0.8 ^a
Phosphorus (%)	0.085 ± 0.013 ^b	0.312 ± 0.036 ^a	0.009 ± 0.001 ^c	0.115 ± 0.003 ^b
Carbon (%)	48.0 ± 0.8 ^a	47.7 ± 0.5 ^a	49.4 ± 0.3 ^a	48.8 ± 0.2 ^a
Nitrogen (%)	2.29 ± 0.03 ^b	0.81 ± 0.03 ^d	1.53 ± 0.08 ^c	2.64 ± 0.12 ^a
C:N (molar)	24.5 ± 0.2 ^c	68.6 ± 2.0 ^a	37.9 ± 1.9 ^b	21.7 ± 0.9 ^c

Values are means ± SE ($n = 3$). Species with different letters are significantly different (one-way ANOVA followed by Tukey HSD test, $p < 0.05$)

Fig. 2 Dry mass remaining, litter toughness remaining and microbial respiration rate on *Alnus glutinosa* (a, e, h), *Quercus robur* (b, f, i), *Acacia melanoxylon* (c, g, j) litter discs and *Acacia dealbata* (d, k) leaflets incubated over 70 days in microcosms simulating conditions of reference and “invaded” streams. Values are means \pm SE ($n = 3$)



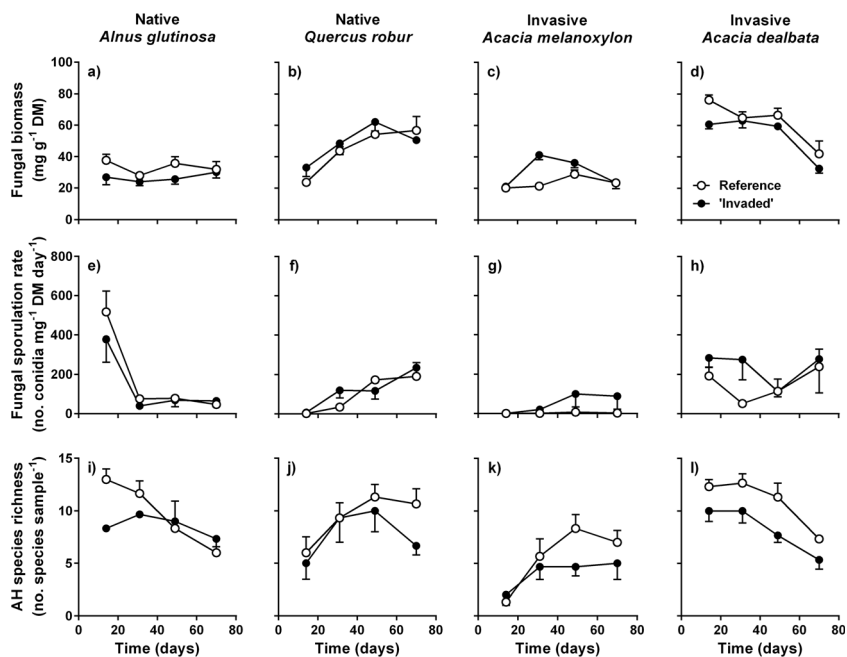
streams, while *Q. robur* and *Ac. melanoxylon* had higher fungal biomass in “invaded” streams (Tukey’s test, $p < 0.001$). Litter species accounted for a higher variance in fungal biomass ($\eta^2 = 54\%$) than the interaction between litter species and stream type ($\eta^2 = 4\%$) (Table S2).

Fungal Sporulation Rate

Maximum sporulation rates by aquatic hyphomycetes varied between 517 and 378 conidia mg^{-1} DM day^{-1} for *Al. glutinosa* litter, 191 and 235 conidia mg^{-1} DM day^{-1} for

Q. robur, 7 and 100 conidia mg^{-1} DM day^{-1} for *Ac. melanoxylon*, and 239 and 284 conidia mg^{-1} DM day^{-1} for *Ac. dealbata*, in reference and “invaded” streams, respectively (Fig. 3e–h). Sporulation rates differed significantly among litter species (three-way univariate PERMANOVA, $p < 0.001$) (Table S3). Generally, *Al. glutinosa*, *Q. robur*, and *Ac. dealbata* litter had higher fungal sporulation rates than *Ac. melanoxylon*, in both reference and “invaded” streams (pair-wise test, $p < 0.050$). Litter species accounted for 18% of the variance in sporulation rates (Table S3). Sporulation rates did neither differ significantly between stream types

Fig. 3 Fungal biomass, fungal sporulation rate and aquatic hyphomycete (AH) species richness associated with *Alnus glutinosa* (a, e, i), *Quercus robur* (b, f, j), *Acacia melanoxylon* (c, g, k) litter discs and *Acacia dealbata* (d, h, l) leaflets incubated over 70 days in microcosms simulating conditions of reference and “invaded” streams. Values are means \pm SE ($n = 3$)



nor there was a significant interaction between litter species and stream type (three-way univariate PERMANOVA, $p < 0.159$ and $p < 0.152$, respectively) (Table S3).

Aquatic Hyphomycete Communities

Maximum aquatic hyphomycete species richness per date varied between 13 and 10 species for *Al. glutinosa* litter, 11 and 10 species for *Q. robur*, 8 and 5 species for *Ac. melanoxylon*, and 13 and 10 for *Ac. dealbata*, in reference and “invaded” streams, respectively (Fig. 3i–l). Species richness differed significantly among litter species and between stream types (three-way ANOVA, $p < 0.001$), with no significant interaction between factors ($p = 0.658$) (Table S2). Aquatic hyphomycete species richness over time was higher in reference than in “invaded” streams and higher for *Al. glutinosa*, *Q. robur*, and *Ac. dealbata* than for *Ac. melanoxylon* litter (Tukey’s test, $p < 0.010$). Litter species accounted for a higher variance in species richness ($\eta^2 = 30\%$) than stream type ($\eta^2 = 6\%$) (Table S2).

Over the entire incubation period, aquatic hyphomycete species richness varied between 23 and 16 species for *Al. glutinosa* litter, 20 and 21 species for *Q. robur*, 17 and 11 species for *Ac. melanoxylon*, and 22 and 20 for *Ac. dealbata*, in reference and “invaded” streams, respectively (Table S4). The aquatic hyphomycete community structure differed significantly among litter species and between stream types (two-way multivariate PERMANOVA, $p < 0.001$). *Articulospora tetracladia* Ingold and *Triscelophorus acuminatus* Nawawi were the most abundant species in *Al. glutinosa*, *Q. robur*, and *Ac. dealbata* litter, in both reference and “invaded” streams (Fig. 4a, b). In *Ac. melanoxylon* litter, *Lunulospora curvula* Ingold was abundant in both stream types, while *A. tetracladia*, *Alatospora pulchella* Marvavová, and *Heliscus lugdunensis* Sacc. & Terry were more abundant in reference streams and *Dimorphospora foliicola* Tubaki was more abundant in “invaded” streams (Fig. 4a, b). Therefore, aquatic hyphomycete communities were distributed in three groups: (i) *Ac. melanoxylon* litter in reference and “invaded” streams, (ii) *Al. glutinosa*, *Q. robur*, and *Ac. dealbata* litter in reference streams, and (iii) *Al. glutinosa*, *Q. robur*, and *Ac. dealbata* litter in “invaded” streams (pair-wise test, $p < 0.050$) (Fig. 5).

Discussion

Invasion of native forests by N-fixing *Acacia* species occurs worldwide [19]. However, information about their effects on the structure and function of freshwater ecosystems is very limited (but see [37–39] for the effects of the invasion of the Fynbos biome by *Acacia mearnsii* De Wild in South African streams). In this study, litter decomposition, microbial

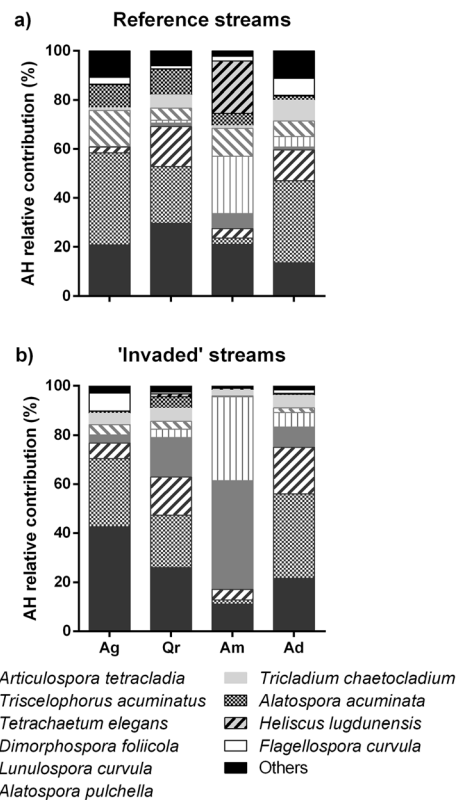


Fig. 4 Relative contribution (% , based on conidial production) of aquatic hyphomycetes (AH) species associated with *Al. glutinosa* (Ag), *Q. robur* (Qr), and *Ac. melanoxylon* (Am) litter discs and *Ac. dealbata* (Ad) leaflets incubated over 70 days in microcosms simulating conditions of reference (a) and “invaded” (b) streams

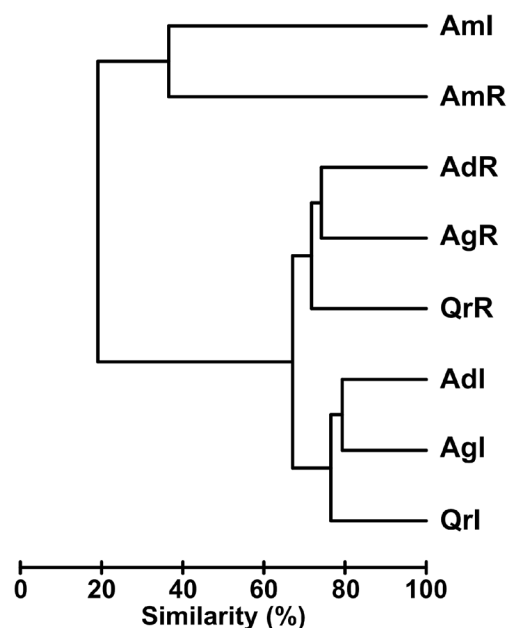


Fig. 5 Cluster dendrogram of aquatic hyphomycete communities associated with *Alnus glutinosa* (Ag), *Quercus robur* (Qr), and *Acacia melanoxylon* (Am) litter discs and *Ac. dealbata* (Ad) leaflets incubated over 70 days in microcosms simulating conditions of reference (R) and “invaded” (I) streams

decomposer activity, and aquatic hyphomycete species richness and community structure differed among litter species, probably due to differences in litter nutritional characteristics, and between stream types most likely due to differences in N concentration in stream water. Our results suggest that invasion of native riparian forests by *Acacia* species can affect litter decomposition and the activity, diversity, and structure of associated microbial decomposers in streams.

Decomposition of *Acacia* Litter Is Slower than that of Native Species

In this study, litter decomposition differed between native and invasive species. Litter decomposition was faster for *Al. glutinosa*, followed by *Q. robur* (native species) and slower for *Ac. melanoxylon* and *Ac. dealbata* (invasive species). This was expected since litter decomposition and microbial decomposer activity strongly depend on the structural and chemical characteristics of litter that fall into streams [40, 41]. Previous studies have shown that soft litter with high nutrient concentrations (e.g., N, P) and low concentrations of structural and secondary compounds (e.g., lignin, polyphenols) generally have higher microbial activity and faster decomposition than litter with high concentrations of recalcitrant compounds [9, 14, 42]. In this study, *Al. glutinosa* litter decomposed faster probably because it was soft, and had a low C:N ratio and a high N concentration, which facilitated microbial colonization and activity. Indeed, the activity of microbial decomposers was higher in *Al. glutinosa* litter when compared with the other species. On the other hand, in *Q. robur* litter, the high polyphenols and low N concentrations were most likely responsible for the limited microbial colonization and activity, and slower decomposition when compared with *Al. glutinosa*. Nevertheless, it was interesting to note that even though *Q. robur* litter was nutrient-poor for microbial decomposers, it decomposed faster than litter of invasive *Acacia* species probably because it was softer than *Ac. melanoxylon* phylloides and had lower lignin concentration than *Ac. dealbata* leaflets. Litter of invasive N-fixing species, *Ac. melanoxylon* and *Ac. dealbata*, also had high N concentrations like *Al. glutinosa*; however, the higher concentration of recalcitrant structural compounds likely contributed to their decreased decomposition. *Acacia melanoxylon* litter decomposed slowly and had lower microbial activity, probably because it had a high lignin concentration and a tough and thick cuticle that acted as a physical barrier for microbial colonization [42]. Previous studies have shown that lignin and toughness are important factors limiting microbial decomposer activity and litter decomposition in streams [11, 42–44]. Therefore, *Ac. melanoxylon* can be considered a poor resource for aquatic microbial decomposers [42]. For *Ac. dealbata*, the reason for slow decomposition was not as clear as leaflets were chemically similar to *Al. glutinosa* litter and were fully colonized by

microbes. One possible explanation is that leaflets probably had a tough and resistant cuticle that limited aquatic hyphomycete enzymatic activity on plant litter tissue, preventing the access of microbes to litter N. Therefore, the higher structural stability of leaflets given by their small size and higher lignin concentration probably allowed fungi to use the inorganic nutrients (e.g., N or P) directly from the stream water to grow mycelia in their surface and sporulate [45]. Indeed, previous studies have shown that higher microbial activity may occur in slow-decomposing substrates [10, 14, 46], probably due to their high structural stability for microbial growth. Additionally, the large surface area:volume ratio of *Ac. dealbata* leaflets may have facilitated its colonization by bacteria, which could have resulted in increased microbial activity but reduced litter decomposition due to the reduced capacity of bacteria to decompose litter [8, 47, 48].

Litter Decomposition Is Faster in “Invaded” than in Reference Streams

In this study, litter decomposition also differed between stream types. Litter decomposition was faster in “invaded” than in reference streams, probably due to the higher NO_3^- -N concentrations found in “invaded” stream water. Previous studies have shown that increases in N concentrations in water can accelerate litter decomposition in streams due to a stimulation of microbial decomposer activity [10, 13, 49, 50]. In our case, differences in decomposer activity between stream types were observed for microbial respiration (*Q. robur*, *Ac. melanoxylon*, and *Ac. dealbata*) and fungal biomass (*Q. robur* and *Ac. melanoxylon*), with higher values in “invaded” than in reference streams. This stimulation of microbial activity in “invaded” streams, although expected, is, nevertheless, impressive given that NO_3^- -N concentrations in “invaded” stream water were still in the oligotrophic range ($< 30 \mu\text{g L}^{-1}$), despite being $\sim 5\times$ higher than in reference streams. Although we did not specifically measure the N concentrations in “invaded” stream water through time, the absence of agricultural activity in the catchment area of “invaded” streams and the absence of other sources of N enrichment suggest that the increased N concentration in “invaded” stream water was most likely due to N leaching from N-rich soils and the decomposition of N-rich litter that falls into streams [25, 28]. Indeed, increased N concentration has also been observed in streams running through forests invaded by other exotic N-fixing species [25–29]. We can therefore anticipate that the invasion of native forests by N-fixing *Acacia* species will most likely increase N concentration in stream water, and that this increase, even if small, will stimulate microbial decomposer activity [10, 49, 51, 52], especially in streams where background N is limiting, and thus, accelerate litter decomposition [10, 53, 54].

Aquatic Hyphomycete Community Structure Differed Among Litter Species and Between Stream Types

Aquatic hyphomycete species richness and community structure differed among litter species, given the differences in litter characteristics [55–57]. Aquatic hyphomycete communities on *Ac. melanoxylon* litter (17 and 11 species in reference and “invaded” streams, respectively) were highly distinct from those on *Al. glutinosa*, *Q. robur*, and *Ac. dealbata*, which were more diverse (20–23 and 16–21 in reference and “invaded” streams, respectively). It was also interesting to note that aquatic hyphomycete species richness and community structure differed between both *Acacia* species. *Acacia dealbata* litter supported higher species richness than *Ac. melanoxylon* and had a community structure very similar to that present on native *Al. glutinosa* and *Q. robur*. Thus, different *Acacia* species can have different effects on aquatic microbial decomposer communities. The invasion of native forests by species with characteristics similar to those of native species may have a weaker effect on stream decomposer communities than the invasion of native forests by species with very different characteristics [2]. However, the replacement of a diverse forest by a monospecific stand may still have strong negative impacts mediated by a reduction in resource diversity [15].

The species richness and structure of aquatic hyphomycete communities also differed between stream types. The total number of aquatic hyphomycete species (across litter species and sampling dates) recorded for reference and “invaded” streams was the same (24 species each), and very similar to the total number of species recorded in this study (25 species). However, the high difference between total and mean aquatic hyphomycete species richness per sampling date suggests high heterogeneity among streams; “invaded” streams had higher heterogeneity than reference streams (i.e., “invaded” streams have different aquatic hyphomycete communities with low number of species, while reference streams have similar communities with high numbers of species resulting in similar overall species richness per stream type). Aquatic hyphomycete communities from reference streams were more diverse than those from “invaded” streams on each sampling date, probably because the first had higher substrate diversity since they flow through diverse deciduous forests, while “invaded” streams flow through forests dominated by *Acacia* species, in some cases forming (almost) monospecific stands [15, 16, 58]. Higher aquatic hyphomycete species richness is expected to occur in streams flowing through diverse forests, due to higher substrate heterogeneity for aquatic hyphomycetes [15, 56]. Nevertheless, aquatic hyphomycete communities in *A. glutinosa*, *Q. robur*, and *Ac. dealbata* litter, from reference and “invaded” streams, were mostly dominated by *A. tetracladia* and *T. acuminatus*. *Articulospora tetracladia* is a species known to have a worldwide distribution [59] and can be found in streams with low and intermediate levels of

nutrient concentrations [50]. Additionally, it was also interesting to note that invasive *Ac. melanoxylon* litter from reference streams supported a higher fungal species richness than litter from “invaded” streams, suggesting that the more diverse communities in reference streams may harbor fungal species with superior enzymatic capabilities (sampling effect mechanism, i.e., higher chance of having specific traits due to higher species richness [60]). In “invaded” streams, the aquatic hyphomycete community from *A. melanoxylon* litter was dominated by *D. foliicola*, a species known to be very abundant in streams with high levels of nutrient concentrations [50, 53]. Although our “invaded” streams were still in the oligotrophic range, the higher NO_3^- -N concentrations in water combined with the recalcitrant characteristics of *A. melanoxylon* stimulated *D. foliicola* conidia production.

Conclusion

Our results suggest that the invasion of native forests by exotic N-fixing *Acacia* species will affect litter decomposition and the activity and structure of associated microbial decomposers. However, the magnitude of the effects may vary with the identity of the *Acacia* species. *Acacia melanoxylon* and *Ac. dealbata* litter decomposed slower than the litter of the native species. However, although decomposer activity and aquatic hyphomycete species richness on *Ac. melanoxylon* litter were lower than on the native species, those on *Ac. dealbata* generally had values in the range of the native species. Also, differences in the structure of aquatic hyphomycete communities were found between native species and *Ac. melanoxylon*, but not *Ac. dealbata*. This suggests that the effects of tree species invasion on stream litter decomposition and microbial communities depend more on the litter characteristics of the species than on the species origin (native/exotic).

Additionally, the differences found between reference and “invaded” streams further suggest that effects of invasion by exotic N-fixing tree species on stream detrital pathways are mediated also by changes in water characteristics (e.g., increase in N concentration), although to a lower extent than by changes in litter characteristics. Overall, invasion of diverse deciduous forests by N-fixing *Acacia* species may affect litter decomposition via multiple pathways (e.g., changes in litter input characteristics, changes in water quality), with the magnitude of the effects depending on the dissimilarity between native and invasive species.

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