Nothing else matters? Tree characteristics have more effects than biogeoclimatic context on microhabitat diversity and occurrence: a nationwide analysis.

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Abstract

Managing forests to preserve biodiversity requires a good knowledge not only of the factors driving its dynamics but also of the structural elements that actually support biodiversity. Tree-related microhabitats (cavities, cracks, conks of fungi) are tree-borne features that are reputed to support specific biodiversity for at least a part of species' life cycles. While several studies have analysed the drivers of microhabitats number and occurrence at the tree scale, they remain limited to a few tree species located in relatively narrow biogeographical ranges. We used a nationwide database of forest reserves where microhabitats were inventoried on more than 22,000 trees. We analysed the effect of tree diameter and living status (alive or dead) on microhabitat number and occurrence per tree, taking into account biogeoclimatic variables and tree genus.

We confirmed that larger trees and dead trees bore more microhabitats than their smaller or living counterparts did; we extended these results to a wider range of tree genera and ecological conditions. Contrary to expectations, the total number of microhabitat types per tree barely varied with tree genus – though we did find slightly higher accumulation levels for broadleaves than for conifers – nor did it vary with elevation or soil pH. We observed the same results for the occurrence of individual microhabitat types. However, accumulation levels with diameter and occurrence on dead trees were higher for microhabitats linked with wood decay processes (e.g. dead branches or woodpecker feeding holes) than for other, epixylic, microhabitats such as epiphytes (ivy, mosses and lichens).

Promoting large living and dead trees of several tree species may be an interesting, and nearly universal, way to favour microhabitats and enhance the substrates needed to support specific biodiversity. In addition, a better understanding of microhabitat drivers and dynamics at the tree scale may help to better define their role as biodiversity indicators for large-scale monitoring.

Introduction

Small natural features are structural habitat elements that have a disproportionately important role for biodiversity related to their actual size [1]. Taking these features into account in biodiversity conservation strategies is a crucial step in science-based decision making [2]. Identifying such structural features in a tri-dimensional forest environment is quite challenging since their number and variety is potentially infinite. Small natural features include, for example, large old trees [3] as well as tree-borne structures. While large old trees are disappearing at the global scale [4], their importance for biodiversity has not yet been fully elucidated, not to mention the peculiar structures they may bear (eg. cracks, cavities, epiphytes), also known as 'tree-related microhabitats' (hereafter 'microhabitats' [5]). Microhabitats have recently aroused the interest of scientists and forest managers alike since these structures can be a substrate for specific forest biodiversity [6], and can ultimately serve as forest biodiversity indicators [5, 7, 8]. Their conservation has hence become an issue in day-to-day forest management, as have large old trees and deadwood [9, 10]. However, our understanding of the drivers and dynamics influencing these microhabitats, notably at the tree scale, remains incomplete [11]. Tree diameter and living status (living vs. dead trees) are key factors for microhabitat diversity at the tree scale [12-14]. Larger trees are likely to bear more microhabitats than smaller ones, as they have experienced more damage, injuries and microhabitat-creating events (e.g. woodpecker excavation, storms, snowfalls). Similarly, gradually decomposing dead trees are likely to bear more microhabitats than living trees and play a role as habitat and food sources for many microhabitat-creating species [15]. Nevertheless, the relationships between microhabitats and tree characteristics have only been demonstrated on a limited number of tree species involving at most a few thousand observations at the tree level (e.g. [11-13]), which have been carried out within a limited biogeographical range (e.g. in Mediterranean forests [16], the French Pyrenees [12] or in Germany [17, 18]). Consequently, it remains to be understood whether the observed relationships between tree characteristics and microhabitats – even though they seem to be relatively consistent across studies – are merely idiosyncratic, notably in terms of magnitude. Large databases making larger-scale analyses possible are

rare (but see [19]), mainly due to a lack of homogeneity in the typologies used to inventory

microhabitats [5] and a lack of forest microhabitat monitoring initiatives. Large-scale data are,

nonetheless, crucial to better understanding the potential variations in the relationships

between microhabitat and biotic (e.g. tree species) or abiotic (e.g. climatic) factors, with a view

to validating microhabitats as potential biodiversity indicators at various scales [7, 8, 18].

We used a nationwide database resulting from standardized monitoring in forest reserves,

where microhabitats have been inventoried since 2005. We analysed the influence of individual

tree diameter and living status on the number and occurrence of microhabitat types at the tree

level. We expected the number and occurrence of microhabitats per tree to increase with

diameter and to be higher on dead than on living trees. We assessed the influence of tree

species and biogeoclimatic variables on these relationships, expecting that microhabitat

dynamics (or accumulation rate per tree) would be tree-species dependent and would vary

with abiotic context (higher accumulation rates in harsher conditions: e.g. at high elevations or

on acidic soils). Ultimately, the aim of this study was to provide forest managers with a better

science-based understanding of microhabitats' role in the forest ecosystem, thus allowing them

to adapt their management to specific local contexts.

Materials and methods

<u>Database structure</u>

We worked with a nationwide database compiled from a monitoring program in French forest

reserves. Since 2005, a systematic permanent plot network has gradually been set-up on a

voluntary basis in forest reserves. The main objectives of this network are (i) to better

understand the natural dynamics of forest ecosystems subjected to varying degrees of

management, (ii) to provide reserve managers with quantitative data on the flux of living and

dead trees at the site scale, and (iii) to ultimately provide guidelines for establishing

management plans. The full database currently includes 107 reserves for a total of 8190 plots

(83180 living and 19615 dead trees, snags or stumps). The forest reserves in the database

actually encompass three broad types of protection status. First, (i) strict forest reserves, where

harvesting has been abandoned for a variable timespan and (ii) special forest reserves, where

management is targeted towards specific biodiversity conservation measures (e.g.

preservation of ponds). These two types are owned and managed by the French National

Forest Service. The third type, nature reserves, on the other hand, where management varies

from abandonment to classic wood production, may be of various ownership types (state, local

authorities, private). It should be noted that no homogeneous data on management intensity

or time since last harvesting could be gathered at the plot level for all the reserves in the

database. However, Vuidot et al. [13] showed that management has a limited effect on

microhabitat number and occurrence at the tree level. We thus assumed that management

differences would not play a significant role at the tree scale and therefore, did not take

management type or intensity into account in our analyses (but see discussion).

<u>Stand structure and microhabitat inventories</u>

On each plot, we combined two sampling methods to characterise forest stand structure [20].

For all living trees with a diameter at breast height (DBH) above 30 cm, we used a fixed angle

plot method to select the individuals comprised within a relascopic angle of 3%. Practically,

this meant that sampling distance was proportional to the apparent DBH of a tree. For example,

a tree with a DBH of 60 cm was included in the sample if it was within 20 m of the centre of the

plot. This particular technique allowed us to better account for larger trees at a small scale. All

other variables were measured on fixed-area plots. Within a fixed 10-m (314 m^2) radius around

the plot centre, we measured the diameter of all living trees and snags (standing dead trees

with a height > 1.30 m) from 7.5 to 30 cm DBH. Within a 20-m radius (1256 m^2), we recorded

all snags with a diameter > 30 cm. Whenever possible, we identified all trees, both alive and

dead, to species level. In the subsequent analyses, we grouped some tree species at the

genus level to have sufficient representation in terms of tree numbers. This resulted in the

following groups: Ash, Beech, Chestnut, Fir, Hornbeam, Larch, Maple (90% sycamore maple),

Oak (80% sessile and pedunculate oaks combined, 15% oaks identified to the genus level

only, 5% other oaks – mainly Mediterranean), Pine (64% Scots pine, 22% mountain pine),

Poplar and Spruce. We assumed that tree genus, rather than species, influenced the

relationships we were studying. Unidentified species were excluded from the analyses.

We visually inspected all selected standing trees for microhabitats and recorded their presence

on each tree. Observers attended a training session and were given a field guide with pictures

to help them better determine microhabitat types and detailed criteria to include in the

inventories. Although inventory methods have recently improved [5, 21], we assumed that the

method we used limited any potential observer effect linked with microhabitat inventories [22].

Different microhabitat typologies were used concomitantly during the inventories and

harmonization has been lacking since 2005. Consequently, we only retained data with a

homogeneous typology. We preferred this solution to grouping microhabitat types to avoid

coarser classification with too much degradation of the original dataset.

<u>Data selection and biogeoclimatic variables extraction</u>

First, we focused on the microhabitat typology that was used for the largest number of plots

and sites (Table 1). This reduced the dataset to 43 sites comprising 3165 plots (Figure 1).

Second, the smallest trees (7.5 ≤ DBH ≤17.5 cm) accounted for 36% of the trees in the

database but were also the least likely to bear microhabitats [12, 13]. We therefore excluded

this category from the dataset to avoid zero-inflation in the subsequent models. Third, previous

studies had shown that tree living status (i.e. living vs. dead trees, see below) is a major driver

of microhabitat occurrence and density [12, 13]. To properly account for this variable in our

statistical models, we excluded all tree species/genera with less than 50 standing dead trees

or snags in the dataset (ie. Ash, Chestnut, Hornbeam, Larch, Maple, Poplar, see Table 2 and

Supplementary Material, Figure S1, for a calculation based on a larger subset of living trees).

The final dataset comprised 2783 plots distributed over 43 sites, for a total of 22307 trees

(20312 living and 1995 dead trees belonging to 11 genera).

In addition, we gathered different biogeoclimatic data from various sources to reflect plot

characteristics:

-   Annual mean temperature (bio1) and precipitation (bio12) from the Worldclim2

database [23];

-   Elevation, aspect and slope from the national digital elevation model (resolution 30 m);

-   Soil plant-bioindicated pH from the National Forest Inventory [24].

<u>Statistical analyses</u>

Following Zuur et al. [25], preliminary data exploration did not reveal any potential variation in

the relationship between microhabitat metrics and any of the biogeoclimatic variables

mentioned above, apart from pH and elevation. Elevation correlated strongly to tree species;

indeed, only Beech and Pine were distributed over the whole elevation gradient while the other

species were elevation-dependent. We therefore kept pH and elevation only in the analyses

described below.

We used DBH, living status (alive vs. dead) and genus (Beech, Fir, Oak, Pine and Spruce) as

explanatory variables and included second and third order interactions between DBH, living

status and genus in the models. We added elevation and pH as covariables, but only included

pH in the second order interactions. Since Beech and Pine were not strongly biased by

elevation, we added elevation in the second order interactions for these two genera in two

separate analysis.

To model the total number of microhabitat types per tree, we used generalised linear mixed

models (GLMMs, library glmmTMB, [26]) with a Poisson error distribution for count data and

plot identity nested within site as a random variable. We also modelled the occurrence of each

microhabitat type, but with a binomial error distribution for binary data. We tested differences

in microhabitat numbers and occurrences between living and dead trees with post-hoc multi-

comparison Tukey tests for a fixed mean DBH (44 cm; function cld, library emmeans [27]).

Dispersion diagnostics revealed under-dispersed model estimations, which may cause a

type II error rate inflation [28]. However, since there was no simple way to account for that in

a frequentist framework, we kept the results while bearing in mind that they were undoubtedly

conservative despite the large number of observations we analysed. In addition, we focused

our interpretations on the magnitude of the results rather than their statistical significance (see

e.g. [29]). We processed all the analyses with the R software v. 3.4.3 [30].

184 Results

<u>Number of microhabitat types per tree</u>

Estimates for all single parameters except for soil pH were significant in the model, while

second and third order interactions were less often significant (see Supplementary Materials,

Table S1). All tree genera except Pine had higher microhabitat numbers on dead than on living

trees. Overall, the difference was the highest for Oak (22% more microhabitats on dead trees,

for a mean DBH of 44 cm, Table 3); the other genera had around 10-15% more microhabitats

on dead trees. Globally, the number of microhabitats per tree increased with tree diameter,

both for living and dead trees (Figure 2). However, the accumulation of microhabitats with

diameter varied with genus: broadleaves (Beech and Oak) had higher accumulation levels

than conifers (Fir, Pine, Spruce); and according to living status: dead versus living trees (except

for Pine; Figure 2, Supplementary Materials, Table S2). These results were generally

consistent with those obtained with the analyses concerning a higher number of genera but for

living trees only (Figure S1). Broadleaves (Ash, Beech, Chestnut, Hornbeam, Maple, Oak,

Poplar) showed higher microhabitat accumulation rates than conifers (Fir, Larch and Spruce).

Only Pine showed accumulation rates comparable to broadleaves (Figure S1).

Number of microhabitats increased significantly with elevation, but not with soil pH. However, higher soil pH had a positive effect on the accumulation of microhabitats with DBH (the second order interaction was significant), mostly on dead trees (Supplementary Materials, Table S1). Still, the effects of elevation and soil pH remained small compared to those of DBH and living status.

For Beech and Pine, the overall results converged with those of the complete model. Soil pH and elevation only had significant effects in the interaction terms (Supplementary Materials: Table S3): increasing soil pH increased microhabitat accumulation with DBH for both species, with a stronger effect for Pine than for Beech. On the other hand, increasing soil pH decreased microhabitat richness on living compared to dead trees. Elevation interacted significantly with living status for Beech only, and almost doubled the difference between living and dead trees; whereas for Pine, the effects were only marginally significant ($p < 0.1$), though high in magnitude.

Occurrence of microhabitat types per tree

Five microhabitats out of twenty generally occurred more frequently on standing deadwood than on living trees, though this was not systematic for all genera or even for living status: trunk cavities (except Fir), woodpecker feeding holes (Figure 3), rot (except Fir), conks of fungi, bark characteristics and crown skeleton (except Fir, Table 3 and Supplementary Materials, Table S4). We observed the strongest differences for woodpecker feeding holes: whatever the genera, they virtually only occurred on standing dead trees (i.e. they were nearly absent from living trees, Figure 3, Table 3). Conversely, injuries, dead branches whatever their size and forks (broadleaves only) occurred more frequently on living trees. Magnitudes for microhabitats more frequent on living trees were around 60% to 90% (Table 3).

For most microhabitats, the probability of occurrence increased with DBH both for living and dead trees, with the remarkable exceptions of canopy cavities, woodpecker cavities and crown

skeletons (Supplementary Materials: Figure S2, Table S4). However, the magnitude of the relation varied with tree genus and living status. For some microhabitat types, the increase in probability of occurrence with DBH was stronger for dead than for living trees, e.g. +35% base and trunk cavities on dead vs. +18% on living Beech, or +23 to +42% for woodpecker feeding holes on dead vs. +0.2 to +3% on living trees (Table S2). Conversely, the increase in probability of occurrence of small and medium dead branches was stronger for living trees (e.g. +53% medium dead branches on living vs. 0.7% on dead Oak) and, to a lesser extent, for mosses on Beech and Fir (+20% and +24% on living trees, vs. +9% and +16% on dead trees, respectively). All other increments with DBH for living trees were smaller, generally below 10%. Note that in some cases, due to the very limited number of occurrences for some microhabitats on certain tree genera, the estimates proved unreliable (huge confidence intervals, e.g. canopy cavities on Oak, Pine and Spruce, Supplementary Materials: Figure S2, Table S4).

Elevation had an overall negative effect on microhabitat occurrence, except for trunk cavities, lichens and forks. Conversely, soil pH tended to have a positive effect on microhabitat occurrence, except for conks of fungi. More interestingly, increasing soil pH had a positive effect on the accumulation of some microhabitats when coupled with DBH (indicated by a significant interaction term), but a negative effect on occurrence on living trees (Supplementary Materials: Table S4). All these significant effects exhibited widely varying levels of magnitude, and in several cases, the estimates were rather imprecise (Supplementary Materials: Figure S2, Table S4).

Discussion

Numerous recent studies in a variety of contexts have shown that the number of microhabitats per tree as well as the occurrence of some microhabitat types increase with tree diameter [11, 14, 16]; the studies also show higher occurrence levels on dead than on living trees [12, 13].

Our nationwide study based on a large database confirmed these relationships and extended

them to a larger range of tree genera under wider biogeographical conditions. Indeed, our

results include five tree genera for both living and dead trees and eleven genera when only

living trees were taken into account (Supplementary Materials: Figure S1).

<u>Dead trees bear more microhabitats than living trees</u>

Standing dead trees contribute significantly to the supply of microhabitats; overall, they bore

10 to 20% more microhabitats than their living counterparts in our dataset. Dead trees often

bear considerably more microhabitats than living trees when individual microhabitat types are

analysed (e.g. woodpecker feeding holes – Figure 3 – or bark characteristics). Once dead,

standing trees are affected by decomposition processes that trigger microhabitat genesis [15].

Standing dead trees also constitute privileged foraging grounds for a number of species [5, 7,

8], including woodpeckers [31, 32]. In particular, insect larvae or ants that live under the bark

of more or less recently dead trees constitute a non-negligible part of some birds' diet [8, 33,

34]. Furthermore, as living trees also bear microhabitats, it seems logical that many of these

would persist when the tree dies and would continue to evolve, or possibly even condition the

presence of other microhabitats linked with the decaying process [15]. For example, injuries

caused by logging, branch break or treefall could begin to rot, then slowly evolve into decay

cavities [5, 35]. These successional changes are likely to explain why these microhabitats

types are more numerous on dead trees. The only exceptions to this global pattern concerned

epiphytes and forks with accumulated organic matter, which both tend to be more numerous

on living trees. Ivy, mosses and lichen are likely to benefit from bark characteristics (e.g. pH,

[36]) occurring only on living trees. Epiphytes, especially slow-growing mosses and lichens,

require a relatively stable substrate to take root and develop [37]. Stability is lost when bark

loosens and falls off during tree senescence, and this could cause epiphytic abundance to

decrease. In a nutshell, decaying processes linked to the tree's death reveal a clear difference

between microhabitats that are linked to decay (i.e. saproxylic microhabitats, sensu [5]) and

those that are not – or less so (i.e. epixylic microhabitats).

Previous studies comparing microhabitat numbers on living and dead trees almost all found

more microhabitats on dead trees (see [17]). However, the degree of difference varies across

studies, from 1.2 times as many microhabitats in Mediterranean forests [16] and twice as many

in five French forests [13] to four times as many on habitat trees in south-western Germany

[38]. Our results ranged from 1.1 to 1.2 times as many microhabitats on dead as on living trees,

which is of a slightly lower order of magnitude than previously reported. This surprising result

may be due to the fact that our study encompassed more species with a lower microhabitat

bearing potential (namely conifers). Yet, even for the same species analysed in previous

studies (e.g. Beech), the levels we observed were lower, and this despite the small effects of

pH and elevation; this finding seems to indicate that the effect is not due to biogeographical

variation.

<u>Number and occurrence of microhabitats increase with tree diameter</u>

We confirmed that both microhabitat number and occurrence increase with tree diameter but,

contrary to expectations ([11-13], but see [14]), tree genus had a limited effect on this

relationship, with only slightly higher microhabitat accumulation levels on broadleaves than on

conifers. Almost all microhabitat types taken individually showed the same increasing trends

with tree DBH, but there were considerable variations in magnitude. Larger (living) trees have

generally lived longer than smaller ones, and are consequently more likely to have suffered

damage during their lifespan due to meteorological events (storms, snowfall), natural hazards

(rockfalls) or use by different tree- and wood-dependent species (woodpeckers, beetles, fungi,

e.g. [13, 39]). Doubling tree diameter (from 50 to 100 cm) roughly doubles the number of tree

microhabitats [13, 17, 18], though some studies have found multiples of up to four [38] or even

five times [12] in certain cases. Again, our results showed magnitudes below the lower end of

this range (the multiplication coefficient ranged from 1.2 to 1.4). This may be because the

largest trees in our dataset were undoubtedly younger than those in the other studies,

especially in studies on near-natural or long-abandoned forests [12, 13]. Indeed, since most of

our sites had been (more or less) recently managed, selective felling may have cause trees

with a given diameter to be younger than their counterparts in primeval forests, where

competition levels may be higher and cause slower growth rates. At the individual microhabitat

scale, dead branches were more likely to occur on large trees than on smaller trees; although

this result seems quite obvious, it had rarely been quantified before. Larger trees have more,

but also larger, branches likely to die from competition with neighbours, especially in

broadleaves [40]. Indeed, Oak and Beech were the genera that showed the highest large dead

branch accumulation rates with diameter in our analyses, while conifers had almost no large

dead branches.

Cavity birds and bats are reputed to prefer larger trees for nesting or roosting [41, 42], since

thicker wood surrounding the cavity provides a better buffered and more stable microclimatic

conditions [43]. However, we did not confirm this relationship; the accumulation rates of

woodpecker cavities with tree diameter was very weak and non-significant. The supposed

relationship between tree diameter and woodpecker cavity occurrence seems hard to prove in

the context of temperate European forests, at least with data from censuses comparable to

ours (see [13] at the tree scale, or [44] at the stand scale); more targeted research focusing on

this specific relationship is probably needed [31, 45]. Our results could also be linked to the

non-linear dynamics [11] of this particular microhabitat. Some cavities in living beech can close

back up when they are no longer used [pers. obs. Y.P.], and trees weakened by cavity digging

can break, e.g. [45]. Other microhabitats, for instance conks of fungi, may also show non-linear

dynamics linked with specific phenology [46]. In our study, the number and occurrence of

microhabitats also increased with diameter in standing dead trees, sometimes at a higher rate

than for living trees. The longer persistence of large dead trees compared to smaller ones [47]

may combine the effects of increased damage due to hazards and the natural decaying

processes described above. This probably explains the higher accumulation levels we

observed in many cases, especially for saproxylic microhabitats (e.g. rot, feeding holes, trunk

cavities). Once again, the only exception to this rule were the epiphytes: their probability of

occurrence tended to increase with tree diameter but very noisily, both for living and dead

trees. For such epiphytic organisms (ivy, mosses and lichens), larger scale processes and

biogeoclimatic context (e.g. soil fertility, precipitation) is probably more important than

individual tree characteristics [48]. This is suggested by the significant and rather strong effects

of pH and elevation in our analyses (Supplementary Materials, Table S4).

<u>Limitations and research perspectives</u>

Contrary to our expectations, we found a limited effect of biogeoclimatic variables on the

relationship between microhabitats, tree diameter and living status. However, some specific

interactions may exist, especially in the case of epiphytes [48], but that could not be evidenced

by our approach. In addition, it was rather difficult to disentangle the effects of tree genus from

those of the biogeoclimatic variables, since the distribution of most tree genera is driven largely

by climate - apart from Beech, and more marginally Pine, which occur over broad bioclimatic

gradients. However, even when we analysed Beech and Pine separately, we did not find any

effect of soil pH or elevation on the number of microhabitats, and only slight effects on

accumulation levels with diameter. These results need to be confirmed by further analyses

with larger, more carefully controlled biogeographical gradients.

Our data from forest reserves potentially reflect a larger anthropogenic gradient than classical

managed forests. Some of the reserves had not been harvested for several decades and

exhibited characteristics of over-mature forests (see e.g. [20], who analysed some of the

reserves included in this paper). On the other hand, their overall structure reflected relatively

recent management abandonment – if any – since the reserves were marked by probable

intensive use or previous harvesting over the past centuries, as is characteristic of western

European forests [49]. This is testified to in the dataset we analysed by the relatively rare

occurrence of dead standing trees, in particular those with a large diameter: standing dead

trees represented a mere 10% of the total dataset and very large individuals (DBH > 67.5cm)

only 1% (Table 2). As a consequence, despite the fact that we worked on an extended

management gradient ranging from managed forests to unmanaged strict reserves, some of

the elements characteristic of old-growth and over-mature forests were still lacking, especially

large dead trees [50]. This truncated our relationships and made them imprecise for the larger

diameter categories. Further research on the last remnant of old-growth primeval forests in

Europe [51, 52] is therefore needed to bridge this gap and better understand microhabitat

dynamics over the whole lifespan of the tree.

Compared to recent developments [5, 21], the microhabitat typology we used (Table 1) seems

rather coarse or imprecise. This may explain why we were not able to confirm some of the

effects mentioned in the literature; different microhabitats from a given group may have

different requirements and dynamics (e.g. cavities dug by the black woodpecker vs. other

woodpecker species). On the other hand, our descriptions allowed us to have enough

occurrences in each type to analyse the combined effects of diameter and genus for almost all

the microhabitat types in the typology. Our approach can be viewed as a compromise between

providing the necessary sample size for statistical analyses and the degree of refinement in

typology. The current developments mentioned above [5] will certainly help to homogenize

data in the near future and to build larger, shared databases on common, comparable grounds.

Despite a training session prior to the inventories, observer effects cannot be totally ruled-out.

Our censuses were mostly performed by non-specialists [22], contrary to the scientific studies

previously published, and this may have led to the relatively low magnitudes observed, with

the hypothesis that detection error is higher on one status (either dead or living trees) or one

type of tree (e.g. small trees, which can be overlooked to the benefit of larger individuals). Such

issues remain to be explored.

Finally, our models assumed – unrealistically as it turns out – that microhabitat number would

increase exponentially with diameter. In fact, recent studies, as well as ecological theory (e.g.

species-area relationship), tend to show a saturated (e.g. logarithmic or sigmoid) relationship

between microhabitats and diameter. Models allowing for different link functions – probably

within a Bayesian framework – will need to be tested to see whether they perform better than

the ones used here (see e.g. [11]).

<u>Implications for forest management and biodiversity conservation</u>

Large old trees are considered keystone small natural features in forest and agro-pastoral

landscapes because of their disproportionate importance for biodiversity relative to their size

[3]. This role for biodiversity is further enhanced by the 'smaller' natural features –

microhabitats – they bear [7]. In our large-scale analysis, we confirmed and extended results

previously observed only locally: most microhabitats occur on large trees, and even more on

dead ones than on living ones. This relationship seems true for several tree genera included

in this analysis, and across a large gradient of ecological conditions, with minor variations in

accumulation rates with soil pH and elevation. As a consequence, conserving and recruiting

large living and dead trees in daily forest management will enhance structural heterogeneity

at the stand scale [6, 53], and favour a variety of tree-borne microhabitats, which could further

help to better conserve specific forest biodiversity [5, 54]. Even though the diameter effect

seems consistent across different conditions, we recommend promoting a variety of large trees

of various species as this may further increase the positive effect on biodiversity [7]. Indeed,

the succession dynamics and formation rate of microhabitats may vary with tree species [11,

13]. The successional patterns and long-term dynamics of microhabitats remain largely

unknown [11] and long-term monitoring at both tree and stand scales are needed to better

understand their dynamics and the underlying processes at play [5]. Ultimately, such

knowledge will provide robust scientific grounds on which to base biodiversity preservation

recommendations for forest managers.

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Figure 1: Location of the study sites

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Figure 2: Relationship between number of microhabitats (N microhabitats) per tree and

Diameter at Breast Height (DBH, cm) by species and living status (living vs. dead standing

trees). Lines represent estimates from generalized mixed effect models with a Poisson error

distribution and plot nested in site as a random effect. Ribbons show the 95% confidence

intervals of the mean. For this representation, pH and elevation were held constant (mean

values in our data set).

Figure 3: Relationship between occurrence (N microhabitats) of woodpecker feeding holes and

Diameter at Breast Height (DBH, cm) by species and living status (living vs. dead standing

trees). Lines represent estimates from generalized mixed effect models with a Binomial error

distribution. Ribbons show the 95% confidence interval of the mean. For this representation,

pH and elevation were held constant. See Supplementary materials, Figure S2, for all

microhabitat types.

Table 1: Microhabitat typology

Microhabitat	Description	Occurrence (%, n=22307)
Base Cavity	Non-woodpecker cavity located at a height < 1.3m, large enough to host small mammals	9.2
Trunk Cavity	Non-woodpecker cavity located at a height comprised between 1.3m and the first main branch	4.5
Canopy cavity	Non-woodpecker cavity located on canopy branches (unhealed)	1.0
Woodpecker cavity	Woodpecker nesting cavity, minimum diameter 2cm	1.4
Crack	Crack in the wood with a width >1cm and deep enough to host bat species	3.1
Woodpecker feeding hole	Feeding hole dug by a woodpecker	4.6
Rot	Presence of wood rot	3.3
Injury	Fresh injury, minimum diameter 10cm.	12.1
Conk of fungi	Conk of a perennial polypore	4.0
Bark characteristic	Bark loosened affecting >50% of the surface of a given part of the tree (base, trunk, canopy)	3.1
Bryophyte (>50)		53.5
Lichen (>50)	Epiphytes with a cover >50% of a given part of the tree (base, trunk, canopy)	31.9
Ivy (>50)		7.9
Small branches (5-10cm)	Dead branches with a diameter comprised between 5 and 10cm and a length > 1m	28.4
Medium branches (10-30cm)	Dead branches with a diameter comprised between 10 and 30cm and a length > 1m	13.3
Large branches (>30cm)	Dead branches with a diameter > 30cm and a length > 1m	1.5
Crown skeleton	Noted when the cumulative number of small, medium and large branches was > 10	2.3
Fork	Fork with suspected presence of organic matter or rainwater	12.8
Broken stem	Broken or dry main stem	7.1

576   Table 2: Distribution of the data by genus and Diameter at Breast Height (DBH) classes: small

577   trees (ST: 17.5 < DBH ≤ 30 cm), medium trees (MT, 30 ≤ DBH < 47.5 cm), large trees (LT,

578   47.5 ≤ DBH < 67.5 cm) and very large trees (VLT, DBH ≥ 67.5 cm). Genera in grey were

579   excluded from the main analyses due to an insufficient number of occurrences of dead trees;

580   in this case, only living trees were analysed (see Supplementary Materials: Figure S1).

	Living trees					Dead trees				
Genus	ST	MT	LT	VLT	Total	ST	MT	LT	VLT	Total
Ash	300	292	93	25	710	25	11	3	0	39
Beech	1743	3382	1811	600	7536	117	213	100	37	467
Chestnut	71	154	87	26	338	42	14	4	3	63
Fir	807	1440	1339	698	4284	126	348	155	54	683
Hornbeam	223	156	30	2	411	8	4	1	0	13
Larch	114	312	243	79	748	6	11	2	0	19
Maple	375	472	140	19	1006	21	10	3	0	34
Oak	1259	1549	1043	925	4776	79	89	38	33	239
Pine	363	783	273	33	1452	83	115	25	5	228
Poplar	66	124	50	18	258	12	11	6	2	31
Spruce	540	850	544	330	2264	87	198	70	26	381
Total	5861	9514	5653	2755	23783	606	1024	407	160	2197

582 Table 3: Percentage of difference between living and dead trees for a mean Diameter at Breast

583 Height (DBH = 44 cm) calculated as [(Microhabitats dead trees – Microhabitats living trees) /

584 (Microhabitats dead trees + Microhabitats living trees)]. An * indicates a significant (p<0.05)

585 difference based on post-hoc Tukey tests for a mean DBH. Values close to 100 correspond to

586 cases where microhabitats were quasi-absent on dead trees (resp. -100 for living trees).

Microhabitats	Beech	Fir	Oak	Pine	Spruce
All	14.9*	12.7*	21.5*	14.8	11.4*
Base cavities	18.6	29.7	4.6	61.3	-32.9
Trunk cavities	41.4*	71.4	49.3*	79.9*	86.8*
Canopy cavities	-44.9	-10.5	10.0	-100.0	100.0
Woodpecker cavities	77.9*	39.7	64.6*	63.7	26.3
Cracks	42.9*	41.4	82.8*	-66.2	54.4
Woodpecker feeding holes	97.5*	98.6*	95.8*	95.6*	97.9*
Rot	45.9*	22.3	90.3*	82.2*	80.3*
Injuries	-67.4*	-82.8*	-62.5*	-74.5*	-89.3*
Conks of fungi	96.1*	98.0*	86.9*	94.1*	96.2*
Bark characteristics	92.1*	94.0*	98.6*	96.9*	98.5*
Moss cover >50%	-18.1*	-37.7*	-56.6*	55.0	6.0
Lichen cover > 50%	-61.1*	-71.9*	-29.1	-32.7	-75.8*
Ivy cover >50%	-25.6	-54.2	-4.5	25.5	-30.9
Small branches	-82.7*	-52.8*	-88.1*	-84.6*	-46.7
Medium branches	-58.8*	81.7*	-59.5*	-48.9	-39.4
Large branches	33.7	42.8	-52.8	54.2	-100.0
Crown skeleton	98.3*	74.6	97.4*	85.3*	91.2*
Forks	-94.3*	-72.4*	-48.9	-82.9	-67.9*
Broken stem	12.1	0.4	-1.6	-38.8	-9.8

Supplementary materials

Table S1: Scaled estimates for number of microhabitat types per tree from a generalised linear mixed model with a Poisson error distribution and plot nested in site as a random effect. DBH: Diameter at Breast Height; SE: standard error of the mean: p = p value; ***p<0.001; **p<0.01; *p<0.05.

Parameter	Estimate	SE	p	
Intercept	0.8198	0.0954	<0.001	***
DBH	0.2265	0.0359	<0.001	***
Fir	-0.3196	0.0482	<0.001	***
Oak	0.2060	0.0502	<0.001	***
Pine	-0.4070	0.0838	<0.001	***
Spruce	-0.4386	0.0558	<0.001	***
Living trees	-0.3004	0.0338	<0.001	***
pH	-0.0170	0.0506	0.7372	ns
Elevation	0.1136	0.0380	0.0028	**
DBH:Fir	-0.0576	0.0469	0.2193	ns
DBH:Oak	-0.0112	0.0474	0.8128	ns
DBH:Pine	-0.1282	0.0886	0.1478	ns
DBH:Spruce	-0.0525	0.0541	0.3318	ns
DBH:Living	-0.0098	0.0368	0.7894	ns
DBH:pH	0.0460	0.0077	<0.001	***
Living:pH	-0.0508	0.0189	0.0072	**
Fir:pH	-0.0362	0.0248	0.1445	ns
Oak:pH	0.0537	0.0221	0.0153	*
Pine:pH	0.0976	0.0350	0.0053	**
Spruce:pH	-0.0311	0.0273	0.2553	ns
Fir:Living	0.0455	0.0491	0.3534	ns
Oak:Living	-0.1354	0.0500	0.0068	**
Pine:Living	0.0017	0.0840	0.9837	ns
Spruce:Living	0.0708	0.0574	0.2173	ns
DBH:Fir:Living	-0.1034	0.0491	0.0352	*
DBH:Oak:Living	-0.0160	0.0484	0.7409	ns
DBH:Pine:Living	0.1964	0.0954	0.0396	*
DBH:Spruce:Living	-0.1154	0.0572	0.0435	*

594   Table S2: Accumulation levels of microhabitats per tree (number of microhabitats and

595   occurrence) for a Diameter at Breast Height (DBH) increment from 50 cm to 100 cm.

	Living trees					Dead trees				
Microhabitats	Beech	Fir	Oak	Pine	Spruce	Beech	Fir	Oak	Pine	Spruce
All	1.173	0.261	0.999	1.641	0.212	1.608	0.919	1.802	0.732	1.006
Base cavities	0.176	0.064	0.052	0.427	0.026	0.345	0.063	0.061	-0.004	0.181
Trunk cavities	0.077	0.013	0.036	0.026	0.05	0.346	0.058	0.282	0.005	0.055
Canopy cavities	0.018	0.002	0.005	0.000	0.000	0.000	0.062	0.002	0.000	0.011
Woodp. cavities	0.005	0.001	0.004	0.209	0.006	0.033	0.027	0.007	0.131	0.023
Cracks	0.028	0.001	0.057	0.007	0.012	0.077	-0.004	0.01	-0.001	-0.003
Woodp. feeding holes	0.032	0.005	0.024	0.002	0.004	0.417	0.305	0.411	0.022	0.233
Rot	0.011	0.003	0.033	0.042	0.001	0.234	0.007	0.267	0.039	-0.019
Injuries	0.043	-0.003	0.043	0.012	-0.004	-0.005	-0.003	0.006	0.012	-0.003
Conks of fungi	0.047	0.022	0.003	0.002	-0.001	0.150	0.240	0.143	-0.01	0.270
Bark characteristics	0.004	0.002	0.013	0.003	0.000	0.021	0.09	0.121	0.014	-0.028
Moss cover >50%	0.196	0.243	-0.032	-0.071	0.246	0.086	0.161	-0.001	0.626	0.277
Lichen cover > 50%	0.063	0.097	0.017	0.174	-0.045	-0.017	0.02	0.005	-0.027	0.026
Ivy cover >50%	0.001	0.002	0.058	-0.002	0.006	0.001	0.000	0.003	0.011	0.000
Small branches	0.200	0.348	0.214	0.147	0.559	-0.003	0.049	-0.015	-0.008	0.007
Medium branches	0.498	0.36	0.526	0.324	0.019	0.015	0.010	0.007	-0.007	0.013
Large branches	0.049	0.003	0.012	0.081	0.000	0.021	0.006	0.000	0.011	0.000
Crown skeleton	0.000	0.001	0.000	0.090	0.005	-0.001	0.000	0.000	0.124	-0.006
Forks	0.279	0.050	0.309	0.152	-0.012	-0.001	0.049	0.001	-0.001	0.048
Broken stem	0.012	-0.021	-0.003	-0.017	-0.01	0.116	0.024	0.052	-0.025	0.052

597 Table S3: Scaled estimates for number of microhabitat types per tree for Beech and Pine from

598 a generalised linear mixed model with a Poisson error distribution and plot nested in site as a

599 random effect. DBH: Diameter at Breast Height; SE: standard error of the mean: p = p value;

600 ***p<0.001; **p<0.01; *p<0.05.

	Beech				Pine			
	Estimate	SE	p		Estimate	SE	p	
Intercept	0.759	0.097	<0.001	***	0.285	0.186	0.126	ns
DBH	0.147	0.034	<0.001	***	0.101	0.046	0.030	*
Living	-0.270	0.033	<0.001	***	-0.441	0.069	<0.001	***
pH	0.032	0.064	0.614	ns	0.173	0.190	0.363	ns
Elevation	-0.030	0.060	0.617	ns	-0.250	0.168	0.137	ns
DBH:Living	0.030	0.034	0.375	ns	0.124	0.051	0.016	*
DBH:pH	0.045	0.010	<0.001	***	0.169	0.067	0.012	*
DBH:Elevation	-0.007	0.009	0.477	ns	-0.105	0.064	0.097	(*)
Living:pH	-0.078	0.033	0.017	*	-0.171	0.155	0.269	ns
Living:Elevation	0.120	0.036	0.001	**	0.302	0.166	0.068	(*)

602 Table S4: Scaled estimates for occurrence of microhabitat types per tree from a generalised linear mixed model with a binomial error distribution
603 and plot nested in site as a random effect. DBH: Diameter at Breast Height; SE: standard error of the mean: p = p value; ***p<0.001; **p<0.01;
604 *p<0.05.

	Base cavities				Trunk cavities				Canopy cavities				Woodpecker cavities				Cracks			
	Estimate	SE	p		Estimate	SE	p		Estimate	SE	p		Estimate	SE	p		Estimate	SE	p	
Intercept	-2.81	0.353	<0.001	***	-2.562	0.266	<0.001	***	-7.499	1.281	<0.001	***	-3.513	0.324	<0.001	***	-3.306	0.324	<0.001	***
DBH	0.85	0.207	<0.001	***	0.824	0.189	<0.001	***	-0.007	1.375	0.996	ns	0.388	0.257	0.131	ns	0.449	0.217	0.039	*
Fir	-0.99	0.344	0.004	**	-1.319	0.304	<0.001	***	-1.223	2.008	0.543	ns	-1.651	0.525	0.002	**	-1.275	0.366	<0.001	***
Oak	-1.077	0.367	0.003	**	-0.409	0.322	0.203	ns	1.389	1.239	0.262	ns	-0.477	0.453	0.292	ns	0.102	0.373	0.784	ns
Pine	-2.019	0.655	0.002	**	-1.256	0.519	0.016	*	-11.291	16.5	0.494	ns	-0.669	0.674	0.321	ns	-3.909	0.879	<0.001	***
Spruce	-1.357	0.343	<0.001	***	-2.06	0.419	<0.001	***	-1.502	2.208	0.496	ns	-2.253	0.647	<0.001	***	-0.842	0.4	0.035	*
Living	-0.395	0.206	0.056	(*)	-0.926	0.183	<0.001	***	0.967	1.21	0.424	ns	-2.11	0.281	<0.001	***	-0.94	0.215	<0.001	***
pH	-0.266	0.286	0.351	ns	-0.256	0.225	0.256	ns	0.969	0.756	0.2	ns	0.256	0.261	0.326	ns	0.125	0.267	0.638	ns
Elevation	0.255	0.219	0.244	ns	0.641	0.176	<0.001	***	-0.011	0.354	0.976	ns	-0.032	0.19	0.867	ns	0.174	0.191	0.361	ns
DBH:Fir	-0.35	0.32	0.275	ns	-0.372	0.28	0.184	ns	2.131	1.571	0.175	ns	0.31	0.429	0.47	ns	-0.641	0.392	0.102	ns
DBH:Oak	-0.298	0.304	0.327	ns	0.125	0.265	0.636	ns	0.663	1.419	0.64	ns	-0.226	0.388	0.56	ns	-0.306	0.321	0.341	ns
DBH:Pine	-0.956	0.625	0.126	ns	-0.682	0.529	0.197	ns	-9.992	13.3	0.454	ns	0.019	0.607	0.975	ns	-1.206	0.745	0.105	ns
DBH:Spruce	0.103	0.321	0.747	ns	-0.259	0.355	0.467	ns	1.65	1.579	0.296	ns	0.326	0.43	0.448	ns	-0.524	0.358	0.143	ns
DBH:Living	-0.188	0.211	0.372	ns	-0.373	0.195	0.055	(*)	0.915	1.378	0.507	ns	0.115	0.27	0.668	ns	-0.028	0.222	0.9	ns
DBH:pH	-0.009	0.054	0.867	ns	0.044	0.062	0.48	ns	-0.146	0.163	0.372	ns	0.129	0.105	0.222	ns	-0.061	0.068	0.368	ns
Living:pH	-0.227	0.359	0.526	ns	-0.881	0.352	0.012	*	-0.757	2.151	0.725	ns	1.266	0.573	0.027	*	0.054	0.403	0.893	ns
Fir:pH	0.301	0.363	0.407	ns	-0.187	0.322	0.561	ns	-1.167	1.231	0.343	ns	0.559	0.468	0.232	ns	-1.461	0.393	<0.001	***
Oak: pH	-1.039	0.644	0.107	ns	-1.285	0.571	0.024	*	10.847	16.4	0.511	ns	0.59	0.674	0.381	ns	2.538	0.748	0.001	**
Pine: pH	1.094	0.348	0.002	**	-1.735	0.616	0.005	**	-10.626	108.9	0.922	ns	1.569	0.649	0.016	*	-0.291	0.451	0.519	ns
Spruce: pH	0.149	0.142	0.293	ns	0.25	0.126	0.048	*	-0.821	0.657	0.211	ns	-0.044	0.196	0.822	ns	-0.434	0.152	0.004	**
Fir:Living	0.07	0.186	0.706	ns	-0.597	0.207	0.004	**	0.601	1.028	0.559	ns	-0.165	0.318	0.603	ns	0.071	0.219	0.747	ns
Oak:Living	0.375	0.177	0.034	*	-0.241	0.187	0.199	ns	-1.067	0.404	0.008	**	-0.345	0.279	0.216	ns	0.053	0.261	0.838	ns
Pine:Living	0.915	0.525	0.081	(*)	0.229	0.41	0.577	ns	0.058	0.9	0.949	ns	0.214	0.387	0.581	ns	1.484	0.509	0.004	**
Spruce:Living	-0.018	0.147	0.902	ns	-0.376	0.297	0.206	ns	-1.108	2.206	0.616	ns	0.295	0.518	0.569	ns	0.111	0.215	0.604	ns
DBH:Fir:Living	0.28	0.334	0.403	ns	0.418	0.32	0.192	ns	-2.472	1.693	0.144	ns	-0.655	0.477	0.17	ns	0.301	0.42	0.474	ns
DBH:Oak:Living	0.155	0.308	0.614	ns	-0.021	0.272	0.938	ns	-0.78	1.423	0.584	ns	0.226	0.405	0.578	ns	0.578	0.331	0.08	(*)
DBH:Pine:Living	1.572	0.684	0.021	*	1.091	0.625	0.081	(*)	9.922	13.357	0.458	ns	0.398	0.666	0.55	ns	1.482	0.747	0.047	*
DBH:Spruce:Living	-0.295	0.333	0.375	ns	0.833	0.438	0.057	(*)	-3.628	128.023	0.977	ns	-0.085	0.463	0.854	ns	0.49	0.394	0.214	ns

Table S4 (continued)

	Woodpecker feeding holes				Rot				Injuries				Conks of fungi				Bark characteristics			
	Estimate	SE	p		Estimate	SE	p		Estimate	SE	p		Estimate	SE	p		Estimate	SE	p	
Intercept	-0.92	0.303	0.002	**	-3.213	0.348	<0.001	***	-4.215	0.404	<0.001	***	-0.534	0.231	0.021	*	-2.726	0.368	<0.001	***
DBH	0.805	0.19	<0.001	***	0.928	0.207	<0.001	***	-0.236	0.342	0.49	ns	0.224	0.15	0.135	ns	0.22	0.197	0.263	ns
Fir	0.352	0.251	0.161	ns	-1.126	0.327	0.001	**	-0.961	0.445	0.031	*	-0.453	0.23	0.048	*	-0.238	0.342	0.487	ns
Oak	0.353	0.274	0.196	ns	1.385	0.321	<0.001	***	-0.258	0.529	0.626	ns	-1.959	0.261	<0.001	***	1.69	0.295	<0.001	***
Pine	-0.98	0.392	0.012	*	-1.083	0.64	0.091	(*)	-1.298	0.54	0.016	*	-2.174	0.563	<0.001	***	-0.094	0.525	0.857	ns
Spruce	-0.573	0.274	0.037	*	-0.389	0.333	0.242	ns	-1.099	0.595	0.065	(*)	-1.195	0.283	<0.001	***	0.595	0.333	0.074	(*)
Living	-4.721	0.23	<0.001	***	-1.018	0.223	<0.001	***	1.701	0.304	<0.001	***	-4.368	0.184	<0.001	***	-3.257	0.222	<0.001	***
pH	-0.107	0.277	0.699	ns	0.523	0.285	0.067	(*)	0.649	0.249	0.009	**	-0.643	0.205	0.002	**	0.673	0.31	0.03	*
Elevation	-0.447	0.207	0.031	*	0.249	0.218	0.255	ns	-0.175	0.145	0.229	ns	-0.512	0.179	0.004	**	-0.41	0.277	0.138	ns
DBH:Fir	-0.281	0.23	0.221	ns	-0.79	0.288	0.006	**	-0.395	0.531	0.457	ns	0.203	0.191	0.287	ns	0.224	0.278	0.421	ns
DBH:Oak	0.227	0.275	0.409	ns	-0.348	0.27	0.198	ns	0.402	0.489	0.412	ns	0.14	0.222	0.529	ns	-0.001	0.268	0.998	ns
DBH:Pine	-0.825	0.368	0.025	*	-0.384	0.559	0.492	ns	0.418	0.544	0.442	ns	-0.364	0.633	0.565	ns	-0.098	0.48	0.838	ns
DBH:Spruce	-0.383	0.261	0.142	ns	-1.395	0.332	<0.001	***	-0.69	0.734	0.347	ns	0.424	0.25	0.09	(*)	-0.493	0.299	0.099	(*)
DBH:Living	<0.001	0.21	0.999	ns	-0.609	0.212	0.004	**	0.52	0.343	0.13	ns	0.367	0.165	0.026	*	0.228	0.209	0.276	ns
DBH:pH	0.162	0.072	0.024	*	0.075	0.065	0.251	ns	0.076	0.034	0.028	*	-0.042	0.063	0.508	ns	0.136	0.088	0.122	ns
Living:pH	-0.665	0.365	0.069	(*)	0.559	0.339	0.099	(*)	0.722	0.446	0.106	ns	-0.564	0.357	0.114	ns	-0.271	0.405	0.504	ns
Fir:pH	0.445	0.3	0.137	ns	-2.106	0.329	<0.001	***	-0.196	0.529	0.711	ns	1.636	0.28	<0.001	***	-1.975	0.312	<0.001	***
Oak: pH	0.786	0.451	0.081	(*)	-1.32	0.7	0.059	(*)	0.248	0.523	0.636	ns	0.809	0.89	0.363	ns	-0.951	0.755	0.208	ns
Pine: pH	-0.037	0.366	0.92	ns	-1.219	0.404	0.003	**	1.253	0.601	0.037	*	0.269	0.45	0.55	ns	-1.72	0.489	<0.001	***
Spruce: pH	0.356	0.141	0.011	*	-0.386	0.15	0.01	*	-0.412	0.146	0.005	**	0.223	0.121	0.066	(*)	-0.983	0.154	<0.001	***
Fir:Living	-0.016	0.227	0.944	ns	-0.356	0.18	0.048	*	-0.565	0.093	<0.001	***	0.513	0.188	0.006	**	0.814	0.321	0.011	*
Oak:Living	0.174	0.219	0.428	ns	-0.305	0.231	0.187	ns	-0.276	0.116	0.018	*	0.275	0.207	0.184	ns	-0.259	0.221	0.242	ns
Pine:Living	-0.204	0.29	0.482	ns	-0.466	0.43	0.278	ns	-0.645	0.157	<0.001	***	0.444	0.361	0.219	ns	0.288	0.386	0.455	ns
Spruce:Living	0.005	0.277	0.986	ns	-0.833	0.227	<0.001	***	-1.081	0.119	<0.001	***	0.164	0.229	0.474	ns	0.193	0.307	0.53	sns
DBH:Fir:Living	-0.232	0.313	0.459	ns	0.737	0.3	0.014	*	0.03	0.535	0.956	ns	-0.226	0.273	0.407	ns	-0.458	0.338	0.175	ns
DBH:Oak:Living	-0.529	0.29	0.068	(*)	0.312	0.282	0.269	ns	-0.409	0.492	0.406	ns	-0.44	0.24	0.066	(*)	-0.204	0.287	0.477	ns
DBH:Pine:Living	0.562	0.478	0.24	ns	0.967	0.727	0.184	ns	-0.671	0.554	0.226	ns	0.464	1.214	0.702	ns	0.048	1.001	0.962	ns
DBH:Spruce:Living	0.058	0.334	0.861	ns	1.1	0.379	0.004	**	0.213	0.739	0.773	ns	-1.409	0.45	0.002	**	-0.036	0.448	0.936	ns

Table S4 (continued)

	Moss (>50%)				Lichen (>50%)				Ivy (>50%)				Small branches				Medium branches			
	Estimate	SE	p		Estimate	SE	p		Estimate	SE	p		Estimate	SE	p		Estimate	SE	p	
Intercept	-0.168	0.618	0.785	ns	-3.5	0.776	<0.001	***	-6.781	0.655	<0.001	***	-4.219	0.429	<0.001	***	-4.441	0.353	<0.001	***
DBH	0.174	0.189	0.358	ns	-0.333	0.27	0.218	ns	0.29	0.407	0.476	ns	-0.212	0.35	0.544	ns	0.359	0.351	0.306	ns
Fir	-1.535	0.256	<0.001	***	0.189	0.318	0.552	ns	-0.889	0.621	0.152	ns	0.774	0.356	0.03	*	1.531	0.341	<0.001	***
Oak	-1.071	0.287	<0.001	***	0.975	0.427	0.022	*	1.084	0.512	0.034	*	0.333	0.432	0.441	ns	1.344	0.363	<0.001	***
Pine	-1.973	0.529	<0.001	***	0.334	0.479	0.485	ns	0.63	0.972	0.517	ns	0.776	0.505	0.124	ns	1.17	0.518	0.024	*
Spruce	-2.116	0.305	<0.001	***	-0.98	0.339	0.004	**	0.36	0.568	0.526	ns	0.341	0.46	0.459	ns	-1.892	1.329	0.155	ns
Living	0.832	0.173	<0.001	***	1.519	0.237	<0.001	***	0.524	0.487	0.283	ns	2.509	0.291	<0.001	***	1.382	0.279	<0.001	***
pH	0.09	0.391	0.819	ns	-0.496	0.453	0.274	ns	1.524	0.492	0.002	**	0.417	0.253	0.099	(*)	0.481	0.224	0.032	*
Elevation	-0.785	0.29	0.007	**	2.353	0.388	<0.001	***	-2.124	0.351	<0.001	***	-0.241	0.158	0.127	ns	-0.237	0.159	0.136	ns
DBH:Fir	0.276	0.273	0.312	ns	0.421	0.343	0.219	ns	-0.046	0.482	0.925	ns	0.615	0.393	0.118	ns	-0.276	0.406	0.496	ns
DBH:Oak	-0.165	0.276	0.55	ns	0.494	0.463	0.286	ns	-0.237	0.449	0.598	ns	-0.822	0.507	0.105	ns	-0.308	0.419	0.462	ns
DBH:Pine	1.284	0.672	0.056	(*)	0.242	0.523	0.643	ns	0.593	1.215	0.625	ns	-0.892	0.542	0.1	ns	-0.459	0.581	0.43	ns
DBH:Spruce	0.586	0.302	0.053	(*)	0.505	0.371	0.173	ns	-0.437	0.534	0.413	ns	0.393	0.458	0.391	ns	0.444	0.799	0.578	ns
DBH:Living	0.362	0.193	0.061	(*)	0.578	0.273	0.034	*	-0.041	0.41	0.921	ns	0.537	0.351	0.127	ns	0.558	0.354	0.115	ns
DBH:pH	0.074	0.042	0.077	(*)	0.046	0.048	0.34	ns	-0.001	0.091	0.991	ns	-0.162	0.032	<0.001	***	-0.022	0.043	0.608	ns
Living:pH	0.21	0.254	0.407	ns	0.499	0.317	0.115	ns	0.692	0.555	0.212	ns	-1.262	0.354	<0.001	***	-3.727	0.385	<0.001	***
Fir:pH	1.853	0.287	<0.001	***	-0.852	0.422	0.043	*	-0.433	0.501	0.388	ns	0.614	0.432	0.155	ns	0.131	0.361	0.718	ns
Oak: pH	-2.15	0.549	<0.001	***	-0.799	0.485	0.1	ns	-1.047	1.018	0.304	ns	0.407	0.496	0.412	ns	-0.239	0.508	0.639	ns
Pine: pH	-0.963	0.305	0.002	**	0.537	0.331	0.104	ns	0.117	0.539	0.828	ns	-1.459	0.472	0.002	**	-0.546	1.373	0.691	ns
Spruce: pH	-0.029	0.11	0.791	ns	-0.383	0.126	0.002	**	-0.523	0.342	0.127	ns	0.005	0.108	0.967	ns	-0.209	0.125	0.094	(*)
Fir:Living	0.377	0.125	0.003	**	-0.718	0.161	<0.001	***	0.802	0.506	0.113	ns	-0.752	0.103	<0.001	***	-0.765	0.163	<0.001	***
Oak:Living	0.172	0.141	0.224	ns	-0.428	0.168	0.011	*	0.488	0.196	0.013	*	-0.074	0.088	0.402	ns	-0.256	0.095	0.007	**
Pine:Living	0.555	0.217	0.01	*	-0.043	0.188	0.819	ns	-0.065	0.397	0.87	ns	-0.302	0.139	0.03	*	0.011	0.176	0.95	ns
Spruce:Living	0.594	0.158	<0.001	***	0.105	0.166	0.525	ns	0.568	0.539	0.292	ns	0.147	0.132	0.268	ns	-0.511	0.258	0.047	*
DBH:Fir:Living	-0.426	0.284	0.134	ns	-0.483	0.351	0.169	ns	0.272	0.496	0.584	ns	-0.201	0.398	0.614	ns	0.456	0.423	0.282	ns
DBH:Oak:Living	-0.42	0.28	0.134	ns	-0.531	0.467	0.255	ns	0.262	0.454	0.564	ns	0.859	0.508	0.091	(*)	0.327	0.422	0.439	ns
DBH:Pine:Living	-2.209	0.739	0.003	**	-0.173	0.568	0.761	ns	-1.185	1.304	0.363	ns	1.251	0.552	0.023	*	0.964	0.597	0.106	ns
DBH:Spruce:Living	-0.704	0.321	0.028	*	-0.895	0.386	0.02	*	0.696	0.549	0.205	ns	0.348	0.471	0.461	ns	-0.337	0.828	0.684	ns

611   Table S4 (continued)

	Large branches				Crown skeleton				Forks				Broken stems			
	Estimate	SE	p		Estimate	SE	p		Estimate	SE	p		Estimate	SE	p	
Intercept	-5.839	0.644	<0.001	***	-5.822	0.933	<0.001	***	-6.104	0.804	<0.001	***	-3.511	0.361	<0.001	***
DBH	0.729	0.499	0.144	ns	0.12	0.304	0.694	ns	-1.374	0.763	0.072	(*)	0.779	0.238	0.001	**
Fir	-1.928	1.504	0.2	ns	-1.784	1.031	0.084	.	0.395	0.775	0.61	ns	0.121	0.31	0.696	ns
Oak	-0.396	0.857	0.644	ns	0.091	0.424	0.83	ns	2.225	0.847	0.009	**	-0.347	0.44	0.43	ns
Pine	1.042	0.924	0.26	ns	0.732	0.637	0.25	ns	0.301	1.077	0.78	ns	-0.867	0.532	0.103	ns
Spruce	-18.493	3608.741	0.996	ns	1.755	0.502	<0.001	***	0.474	0.81	0.559	ns	-0.331	0.355	0.351	ns
Living	-0.703	0.593	0.236	ns	-4.754	0.512	<0.001	***	3.597	0.731	<0.001	***	-0.25	0.266	0.347	ns
pH	0.032	0.403	0.937	ns	-0.394	0.634	0.534	ns	-0.209	0.299	0.485	ns	0.688	0.251	0.006	**
Elevation	-0.214	0.277	0.438	ns	-0.784	0.304	0.01	*	1.543	0.198	<0.001	***	0.278	0.165	0.092	(*)
DBH:Fir	0.314	0.947	0.74	ns	-0.539	0.839	0.52	ns	2.116	0.8	0.008	**	-0.551	0.286	0.054	(*)
DBH:Oak	-0.683	0.742	0.357	ns	-0.05	0.349	0.887	ns	1.43	0.865	0.098	(*)	-0.135	0.331	0.682	ns
DBH:Pine	0.768	0.872	0.378	ns	0.666	0.506	0.188	ns	1.042	1.111	0.348	ns	-1.74	0.512	0.001	**
DBH:Spruce	-1.007	4451.258	1	ns	-0.904	0.495	0.068	(*)	2	0.804	0.013	*	-0.405	0.313	0.195	ns
DBH:Living	0.505	0.51	0.322	ns	-0.049	0.425	0.908	ns	2.028	0.764	0.008	**	-0.53	0.242	0.028	*
DBH:pH	-0.086	0.112	0.443	ns	0.372	0.153	0.015	*	0.138	0.038	<0.001	***	0.09	0.047	0.056	(*)
Living:pH	-0.211	1.74	0.903	ns	2.824	0.933	0.002	**	-1.749	0.777	0.024	*	0.242	0.314	0.442	ns
Fir:pH	1.881	0.863	0.029	*	0.409	0.576	0.478	ns	-2.486	0.845	0.003	**	0.282	0.44	0.521	ns
Oak: pH	-0.516	0.938	0.582	ns	2.212	0.729	0.002	**	-1.196	1.077	0.267	ns	1.084	0.523	0.038	*
Pine: pH	11.885	3608.739	0.997	ns	1.664	0.669	0.013	*	-1.927	0.813	0.018	*	0.451	0.364	0.216	ns
Spruce: pH	-0.01	0.312	0.975	ns	0.372	0.266	0.161	ns	-0.201	0.166	0.226	ns	-0.361	0.111	0.001	**
Fir:Living	0.169	0.728	0.816	ns	1.987	1.066	0.062	(*)	0.682	0.102	<0.001	***	-0.354	0.121	0.003	**
Oak:Living	-0.234	0.278	0.4	ns	0.181	0.576	0.753	ns	0.05	0.136	0.712	ns	0.186	0.203	0.359	ns
Pine:Living	-0.338	0.442	0.445	ns	1.75	0.679	0.01	*	0.319	0.178	0.073	(*)	-0.126	0.188	0.504	ns
Spruce:Living	2.271	6.236	0.716	ns	0.845	0.475	0.075	(*)	-0.295	0.138	0.032	*	-0.325	0.17	0.056	(*)
DBH:Fir:Living	-0.726	1.054	0.491	ns	1.364	0.902	0.131	ns	-2.664	0.803	0.001	**	-0.43	0.297	0.147	ns
DBH:Oak:Living	0.353	0.75	0.638	ns	0.645	0.471	0.171	ns	-1.516	0.866	0.08	(*)	-0.246	0.342	0.471	ns
DBH:Pine:Living	-0.34	0.955	0.722	ns	0.31	0.567	0.584	ns	-1.093	1.121	0.329	ns	0.753	0.532	0.157	ns
DBH:Spruce:Living	2.124	4451.258	1	ns	1.561	0.615	0.011	*	-2.744	0.808	0.001	**	-0.293	0.327	0.371	ns

Figure S1: Relationship between total number of microhabitats per tree (N microhabitats) and Diameter at

Breast Height (DBH, cm) by genus and living status (living vs. dead standing trees). Lines represent estimates

from generalized mixed effect models with a Poisson error distribution. Ribbons show the 95% confidence

interval of the mean. For this representation, pH and elevation were held constant.

Figure S2: Relationship between occurrence of microhabitats per tree and Diameter at Breast Height (DBH,

cm) by species and living status (living vs. dead standing trees). Lines represent estimates from generalized

mixed effect models with a Binomial error distribution. Ribbons show the 95% confidence interval of the mean.

For the representation, pH and elevation were held constant.