

Habitat specificity and diversity of tree species in an African wet tropical forest

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Abstract Niche differentiation with respect to habitat has been hypothesized to shape patterns of diversity and species distributions in plant communities. African forests have been reported to be relatively less diverse compared to highly diversified regions of the Amazonian or Southeast Asian forests, and might be expected to have less niche differentiation. We examined patterns of structural and floristic differences

among five topographically defined habitats for 494 species with stems ≥ 1 cm dbh in a 50-ha plot in Korup National Park, Cameroon. In addition, we tested for species–habitat associations for 272 species (with more than 50 individuals in the plot) using Torus translation randomization tests. Tree density and basal area were lowest in areas with negative convexity, which contained streams or were inundated during rainy periods and highest in moist well-drained habitats. Species composition and diversity varied along the topographical gradient from low flat to ridge top habitats. The low depression and low flat habitats were characterized by high diversity and similar species composition, relative to slopes, high gullies and ridge tops. Sixty-three percent of the species evaluated showed significant positive associations with at least one of the five habitat types. The majority of associations were with low depressions (75 species) and the fewest with ridge tops (8 species). The large number of species–habitat associations and the pronounced contrast between low (valley) and elevated (ridgetop) habitats in the Korup plot shows that niche differentiation with respect to edaphic variables (e.g., soil moisture, nutrients) contributes to local scale tree species distributions and to the maintenance of diversity in African forests.

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Introduction

Understanding the processes that generated and continue to maintain high levels of species diversity in tropical forests is a central challenge to community ecology (Hubbell and Foster 1986; Gentry 1988; Hubbell 2001; Wright 2002; Condit et al. 2005). A long-standing hypothesis for the maintenance of diversity within species-rich forests is that resource-based niche differentiation and other mechanisms of habitat specificity allow large numbers of species to coexist within communities (Ricklefs 1977; Webb and Peart 2000; Harms et al. 2001; Hardy and Sonké 2004; Potts et al. 2004; Valencia et al. 2004; Gunatilleke et al. 2006). Species distribution patterns that are biased with respect to environmental variables would support this hypothesis (Gartlan et al. 1986; Newbery et al. 1988; Clark et al. 1999; Webb and Peart 2000; Phillips et al. 2003; Hall et al. 2004; Paoli et al. 2006).

Topographic features greatly influence local conditions, especially soil processes and hydrology, that may in turn influence the demographic processes of growth, mortality, and recruitment (Gartlan et al. 1986; Clark et al. 1999; Daws et al. 2002; Miyamoto et al. 2003). Variation in demographic performance among habitats can translate into species' associations with their preferred habitats (Comita and Engelbrecht 2009; Russo et al. 2008; Yamada et al. 2007). Thus, resolving the role of habitat partitioning in the maintenance of high species diversity in rainforests will depend in part on detecting species–habitat associations (Webb and Peart 2000; Harms et al. 2001; Hall et al. 2004; Potts et al. 2004). Differences among species in their habitat associations, coupled with habitat heterogeneity, will contribute to the maintenance of high diversity by allowing species to coexist by specializing on different habitats. In addition, individual species–habitat associations will shape spatial patterns of forest structure, species composition and diversity, as habitats considered more conducive will serve as concentration areas for a higher number of species (Potts et al. 2004).

Comparative studies of the main tropical forest ecosystems show that African rainforests have relatively poor α -diversity compared to that of the highest diversity regions of Asia and the Americas (Parmentier et al. 2007). In contrast to this overall pattern of diversity, current understanding of the local-scale community-assembly mechanisms for tropical African

tree communities is very limited and complicated by previous sampling designs. For example, inventories based on 1-ha plots spread across a wide area capture fewer than half of the local species, with many represented only by a single individual (Hall and Swaine 1981). Furthermore, most of these inventories focus on large trees with dbh \geq 10 cm (Hall and Swaine 1981; Gartlan et al. 1986; Newbery et al. 1986; Hardy and Sonké 2004) and in some cases only include selected taxa (Hall et al. 2004). These small plots limit the identification of habitats at scales that could provide meaningful inferences on plant populations and also preclude comparisons of degrees of habitat specificity with other tropical forests.

We established a 50-ha forest dynamics research plot (FDP) in southwestern Cameroon, with the largest single plot dataset from the African continent, to document the abundance of all tree species and evaluate species–habitat associations that reflect niche partitioning as a driving force for species coexistence. The plot includes varied terrain to assess the extent to which tree species in the community have biased distributions with respect to environmental variation defined by topographic habitat types. With the first enumeration of trees completed, we ask in this article the following questions: (i) To what degree do habitats differ in forest structure and species composition? (ii) Are pockets of high or low plant diversity associated with particular habitat types? (iii) To what extent are species' distributions associated with habitat types defined by topographic features? (iv) Are habitats with many associated species the most diverse?

Materials and methods

Study area

Korup National Park (5°10' N, 8°70' E; Fig. 1) is located in southwestern Cameroon and lies within the Guineo-Congolian forest of tropical Africa (White 1983), in a species-rich area that has been identified as a former Pleistocene refugium (Maley 1987). It constitutes one of the last remnants of the Atlantic coastal forest, i.e., *la forêt biafriéenne* (Letouzey 1985), in an area that has been identified as a biodiversity hotspot, combining both high plant diversity with a high rate of deforestation (Küper et al. 2004).

The climate of the southern part of Korup is strongly seasonal, with a single distinct dry season (average monthly rainfall < 100 mm) from December to February, followed by a long and intense wet season, with heaviest rainfall in August (Newbery et al. 1998; Chuyong et al. 2004a). Over a 17-year period from 1984 to 2001, the southern part of the park received an annual average of 5040 mm of rain (Newbery et al. 2004). The average monthly temperature ranges between 25.1 and 27.6°C and solar radiation ranges from 199 to 248 W/m². The soils are derived from highly weathered syenite and other igneous rocks; because of the high rainfall they are skeletal and sandy at the surface, highly leached, and poor in nutrients (Newbery et al. 1998; Chuyong et al. 2002).

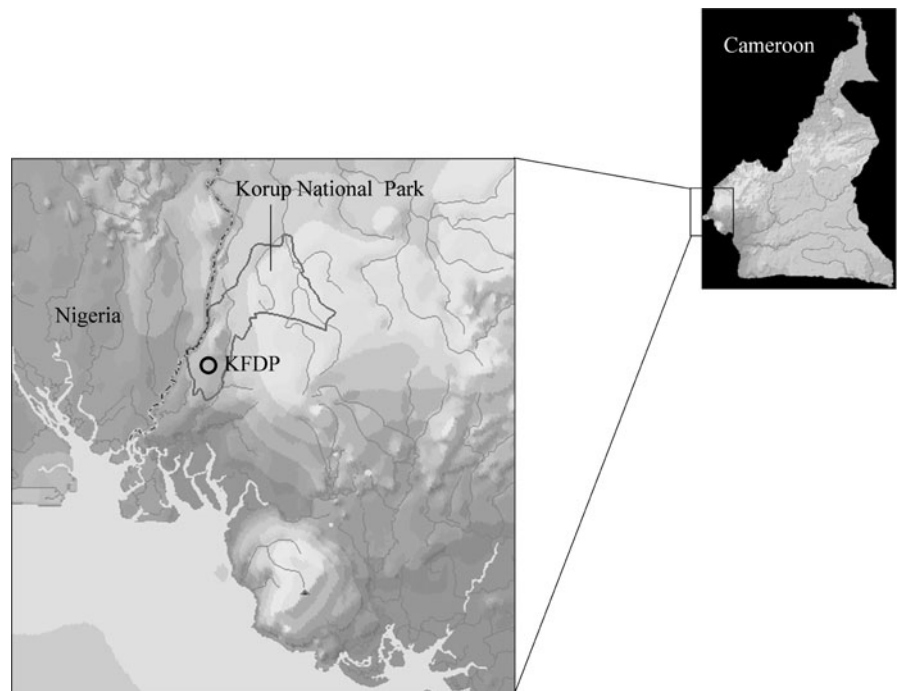
The 50-ha Korup FDP is located in the southern part of Korup National Park and covers an area with diverse terrain (Fig. 1). The plot is 1000 m × 500 m, with the longer axis running north–south and its elevation ranges from 150 to 240 m asl. The southern part of the plot is fairly flat, with a valley bottom that contains a permanent stream flowing westward, while the northern part is steeper, with distinct gullies and large boulders. Further description of the Korup FDP can be found in Thomas et al. (2003), Chuyong et al. (2004b), and Kenfack et al. (2007).

Plot enumeration

The first enumeration of the Korup FDP was carried out from November 1996 to June 1999, following the Center for Tropical Forest Science (CTFS) standardized methods described in Condit (1998). Initially, the 50-ha plot was surveyed and demarcated into 1,250 20 × 20 m quadrats. Elevation was measured to the nearest 0.1 m at each corner post and the value added to the absolute elevation got from the published survey map of the area. Standard geostatistical kriging procedures were then used to produce the topographic map based on a 5-m grid spanning the entire plot (see Fig. 2; kriged points at 5-m intervals were used only to produce the map for visualization purposes—all tests were conducted with the 20-m interval elevation values).

All trees ≥ 1 cm in stem diameter in each quadrat were measured, tagged with unique identification numbers, mapped to the nearest 50 cm, and sorted into morphospecies. Diameter was measured at breast height (DBH, i.e., 1.3 m above the ground) except for irregular stems (swollen and buttressed stems), in which case the point of measurement was just below the swelling, or 0.5 m above buttresses and stilt roots. Most stems were measured at breast height, so we use

Fig. 1 Map of Cameroon and the Forest Dynamics Plot location in Korup National Park



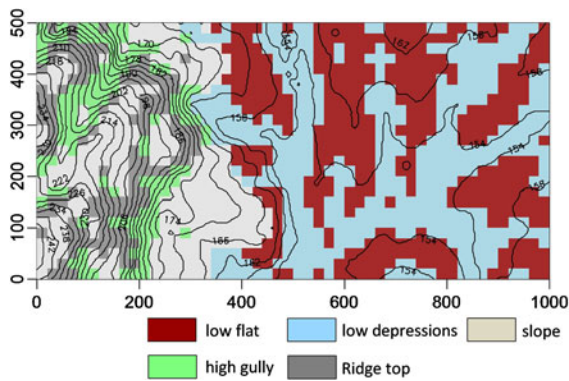


Fig. 2 Topographic map of the 50-ha Korup Forest Dynamics Plot with 2-m contour intervals. The five habitats recognized in this study are indicated: low depressions; low flat; high gullies; slopes and ridge top

DBH to refer to all diameter measurements in this article. The stems were mapped relative to the 20×20 m grid in the field and their respective xy -coordinates were obtained by digitizing these mapped points on a Summersketch[®] tablet (Thomas et al. 2003).

Herbarium specimens were collected for every morphospecies. Taxonomic identification was conducted using regional floras and by matching specimens at SCA, K, MO, and YA (herbarium abbreviations following Holmgren et al. 1990). Specimens were deposited at the respective herbaria where matchings were done.

Close to 80% of the species have been confidently identified and some new to science have been described (see Sonké et al. 2002; Kenfack et al. 2003, 2004, 2006). Family classifications followed the Angiosperm Phylogeny Group (APG II, 2003). Species were grouped into four life-forms defined by their maximum attainable heights as follows; treelets/small trees (<10 m), understorey (10–20 m), lower canopy (20–30 m), and upper canopy (>30 m). Maximum attainable heights were obtained from the literature supplemented by field observations (Thomas et al. 2003).

Habitat classification

Each 20×20 m quadrat was assigned to a habitat category based on its topographic attributes, elevation, slope and convexity, using the methods described in Yamakura et al. (1995), Harms et al. (2001), and

Valencia et al. (2004). For each quadrat, elevation was calculated as the mean of the elevation at its four corners. Slope was the mean angular deviation from the horizontal of each of the four triangular planes formed by connecting three corners at a time. Convexity was calculated as the difference between the mean elevation of the focal quadrat and the mean elevation of the eight surrounding quadrats. For edge quadrats, convexity was calculated as the difference between the elevation of the focal quadrat center and the mean elevation of the four corners. The transition from valley to slope coincides with the uppermost flood level of the stream that flows through the plot and was defined as 165 m, based on observations of high water levels. This elevation is convenient since it is close to the median elevation of the plot. Using thresholds of 165 m for mean elevation, 15° for slope (the median) and zero for convexity, the 1,250 quadrats were classified into five topographic habitats (Fig. 2) as follows: low depression (mean elevation < 165 m, slope < 15° and convexity < 0); low flat (mean elevation < 165 m, slope < 15° & convexity \geq 0); high gully (mean elevation \geq 165 m, slope \geq 15° , and convexity < 0); Slopes (mean elevation \geq 165 m, slope \geq 15° and convexity \geq 0); and Ridge top (mean elevation \geq 165 m and slope < 15°).

Structural and floristic differences among habitats

To compare forest structure among habitats, stem density, and mean basal area were calculated in each 20×20 m quadrat and then the mean and standard deviation were calculated over quadrats assigned to each habitat type. To compare diversity among habitat types, we calculated species richness as the total number of species found in each habitat, as well as the mean number of species per 20×20 m quadrat. To control for differences in stem density and area among habitat types, we also calculated Fisher's α (following the routine given in Condit 1998) for each 20×20 m quadrat and obtained the mean for each habitat type, as well as the overall Fisher's α for each habitat type (i.e., all quadrats combined). Confidence limits were calculated based on variances across quadrats within the different habitat categories.

Pairwise comparisons of species stem density between habitats were used to assess inter-habitat similarity. A straight line through the origin with slope = 1 would signify that all species had consistent

densities in the two habitats. We fitted no-intercept linear regressions to the log-transformed densities and used the corresponding r^2 values as similarity indices. The mean Sørensen similarity index “with cover” (Valencia et al. 2004) was also calculated for pairs of 20×20 m quadrats in habitats i and j for within and between-habitat comparisons in species composition. Between-habitat means were standardized as in Valencia et al. (2004), but without correction for geographic distance using:

$$\frac{100 \times \text{SOR}_{ij}}{0.5(\text{SOR}_{ii} + \text{SOR}_{jj})}$$

where SOR_{ij} is the mean Sørensen similarity index between quadrat pairs (one in habitat i and the other in habitat j). Since the most abundant species are often used to define forest composition (e.g., Valencia et al. 2004), the top 10 ranking species in density in each habitat were considered as dominants and used in assessing compositional differences.

Species–habitat associations

One of our primary objectives was to determine the prevalence of topographic habitat association in tree species in the Korup plot. The challenge to determining how many species and to what extent they show habitat associations is the inherent spatial autocorrelation that occurs both in species distributions and in topography, which is often characterized by large areas of similar features such as large flat areas, or slopes up to a ridge top (Condit et al. 2000; Webb and Peart 2000; Harms et al. 2001; Philips et al. 2003). Therefore, many ordinary parametric and nonparametric analyses are not appropriate and can lead to spurious associations when they assume independence among non-independent sample units (Legendre 1993). Therefore, we adopted torus-translation tests (Harms et al. 2001; Hall et al. 2004; Gunatilleke et al. 2006) to assess patterns of association between trees and our topographically defined habitats.

Our torus-translation tests incorporated the spatial structure of the trees and most of the spatial autocorrelation of the topographically defined habitats. Each test required translating the habitat map in four cardinal directions, moving the entire habitat map one column or row of 20×20 m quadrats at a time. The positions of the individual stems were not altered as the habitats were shifted, but the character

of the habitat assigned to a given stem changed with the shifts of the habitat map. The number of individuals of each species was counted in each quadrat for each translation of the habitat map, and the relative density for each focal species in each focal habitat was calculated as the density of the focal species divided by the density of all species combined in the focal habitat, i.e., the proportion of all stems in a given habitat represented by the focal species. Each of the 1,249 unique translations of the habitat map (not including the 0,0 translation, i.e., the true habitat map) provided a new estimate of the expected relative density of each species in each habitat if species distributions had appeared on the landscape at random with respect to habitats (Harms et al. 2001).

The observed association of a species with a given habitat (on the true habitat map) was compared to the frequency distribution of expected values. In torus translation tests with a significance level of $\alpha = 0.05$, significant associations are those in which the observed level of association (measured by relative density) are ≤ 2.5 or $\geq 97.5\%$ of the expected values. However here, we only consider positive associations, and therefore our tests are essentially one-tailed tests with $\alpha = 0.025$. Rare species with < 1 individual/ha were excluded from the analysis, leaving a total of 272 species for which we tested for significant positive association with the five habitat types.

Results

Habitat differences in stand structure, species diversity and composition

The low depressions had the lowest stem density and high gullies had the lowest basal area, whereas the low flat habitat had the highest stem density and slopes had the highest basal area (Table 1). High stem density in the low flat habitat was mostly due to small sized individuals. High basal area on the slopes matched the higher proportion of large stemmed individuals (dbh > 60 cm) in that habitat.

Species richness varied among habitat types and declined from the lower part of the plot to the ridge top (Table 1). Based on Fisher's α , the low flat habitat was the most diverse and the high gullies the least diverse (Table 1). Three significant diversity groups

Table 1 Density, basal area and diversity of the different habitats within the Korup forest dynamics plot

Habitat	Low depressions	Low flat	High gullies	Slopes	Ridge top
Area (ha)	16.3	15.2	4.3	4.7	9.5
Density (individuals/ha)					
All individuals	5949	7817	5527	6093	6407
1–10 cm	5502	7334	5007	5508	5881
10–30 cm	371	402	418	486	438
30–60 cm	67	70	93	84	77
>60 cm	9	11	9	15	11
Basal area/ha					
All individuals	27.8	33.2	31.6	35.8	32.3
Species richness					
Entire habitat	424	423	346	344	383
Mean/quadrat	67.3 ^b	75.3 ^a	57.9 ^c	64.6 ^b	62.2 ^b
SD/quadrat	12.3	10.8	11.9	17.7	13.1
Fisher's α					
Entire habitat	56.8	55.0	57.1	54.7	54.4
Mean/quadrat	32.09 ^a	32.03 ^a	25.92 ^c	29.22 ^b	26.59 ^c
SD/quadrat	7.18	6.04	7.08	8.38	6.67

Species number and Fisher's α per quadrat were calculated as the mean (± 1 SD) of 20×20 m quadrats. Means with different superscript letters are significantly different

were obtained in the following decreasing order: low flat and low depression > slopes > high gully and ridge tops.

In terms of species composition, low depression and low flat areas contained similar densities of all species. This was evident in the high r^2 value (0.80) obtained from the pairwise regression of species density between the two habitats. High gullies, slopes and ridge tops were also very similar; with r^2 values ranging between 0.76 and 0.86 (see Table 2). The low habitats (low depression and low flat) differed in densities of all species relative to the high habitats (high gullies, slopes and ridge top), as indicated by low r^2 values for pairwise habitat comparisons of density (Table 2). This pattern was also evident in the disparity of the Sørensen similarity indices between the low and high habitats. The low depression and low flat were relatively similar to each other, as were high gullies, slopes and ridge tops (Table 2). The greatest contrast was between low flat areas and high gullies.

Dominant species were inconsistent across the different habitats (Table 3). For example, *Phyllobotryon spathulatum*, the dominant species in the low flat with 921 individuals/ha, had 144 individuals/ha and was ranked 8th in the high gully habitat; *Rinorea gabunensis* was dominant on the slopes with 383 individuals/ha compared to only 3 individuals/ha in the low flat habitat (ranked 177). Even so, similarities

in dominant species occurred between low depression and low flat habitats. Overall, all of the 10 forest-wide dominant species had densities >50 individuals/ha in each habitat, except *Rinorea lepidobotrys*, which had low densities in the high gullies and slopes (Table 3).

Species–habitat associations

Of the 272 species with >50 individuals in the plot, 172 (63%) showed significant positive associations with at least one of the five habitat types (Table 4). The majority of associations were with low depressions (75 species) and the fewest with the ridge tops (8 species). Only 8 species showed multiple positive associations (even the pair of habitats with the most species positively associated with both, i.e., high gullies and slopes, had only five positively associated species in common). The 172 species that showed significant positive associations represented 63 upper canopy, 46 sub-canopy, 48 understorey, and 15 treelets species, which represent 84, 55, 43, and 7% of species in each category, respectively. Selected distribution maps illustrate some of these patterns of association (Fig. 3). For example, *Beilschmiedia* sp. showed strong association to the low depressions (Fig. 3a) *Cola semecarpophylla* and *Lecomtedoxa klaineana* (Fig. 3b) to low flat areas (Fig. 3). *Drypetes staudtii*

Table 2 Comparison of abundance of individual species between habitats using 272 species with ≥ 50 individuals within the entire Korup Forest Dynamics Plot

Habitat comparison	Number of species				Compositional similarity	
	Obligate	Similar	Moderately different	Very different	r^2	Standardised Sørensen index
Low depression vs. low flat	3	135	119	15	0.80	96.8
Low depression vs. high gullies	30	30	83	129	0.17	75.5
Low depression vs. slopes	30	36	96	110	0.11	73.8
Low depression vs. ridge top	15	42	140	75	0.32	83.4
Low flat vs. high gullies	31	27	79	135	0.20	73.7
Low flat vs. Slopes	31	37	127	114	0.19	74.5
Low flat vs. ridge top	16	42	135	79	0.40	84.2
High gullies vs. slopes	38	98	127	9	0.86	97.1
High gullies vs. ridge top	30	35	172	35	0.79	95.6
Slopes vs. ridge top	31	51	150	29	0.76	95.8

“Obligate” refers to species that occurred in only one of the habitats; “Similar” refers to species with density difference < 1.5 -fold between a pair of habitats; “Moderately different” indicates density difference ≥ 1.5 -fold but < 5 -fold; and “Very different” indicates species with ≥ 5 -fold difference in density between the two habitats. The r^2 values are from regressions of all species’ log-density across two habitats (including all 473 species). Standardized similarity is expressed as a percentage of within habitat similarity based on Sørensen’s index

showed strong association to both high gullies and slope. *Hymenostegia afzelii* appears to be a generalist in that it is relatively evenly distributed among habitats (Fig. 3C). Significant habitat association was not restricted to dominant species with relatively high densities in their preferred habitat, but included rarer species as well. For example, *Homalium sarcopetalum* and *Rauvolfia mannii* were significantly associated to the low depression but had densities < 10 individuals ha^{-1} in that habitat.

Discussion

Forest structure, composition and diversity across topographically defined habitats

Our results reveal clear differences in forest structure, diversity, and species composition among the five topographic habitat types in the Korup forest dynamics plot. Tree density and basal area were lowest in habitats with negative convexity values (low depressions and high gullies, respectively). Such areas contain streams or are likely to be inundated during rainy periods, which reduces the area available for tree recruitment and may also lower growth and survival, resulting in fewer, smaller trees. In contrast,

the habitats with the highest tree density and basal area (low flat and slopes, respectively) are both well-drained, but relatively moist areas, suggesting that such habitats are better for overall tree performance than inundated or dry habitats. This is consistent with topography-related patterns of forest structure in central Panama, where tree densities were found to be lowest in inundated (swamp) and streamside habitats and highest in low plateau and slope habitats (Harms et al. 2001).

Interestingly, patterns of diversity and composition among habitats differed from those of forest structure. Diversity per quadrat was highest in low-lying habitats, despite the low tree density in low depressions, and decreased from slopes to high gully and ridge top habitats. Low depressions and low flat habitats also had similar species composition, which differed from the composition in steeper and elevated habitats. These trends in species composition and diversity at Korup follow the ridge-valley contrast that has been reported for other tropical forest sites in Ecuador (Valencia et al. 2004) and Sri Lanka (Gunatilleke et al. 2006). Despite clear compositional differences among those habitat groups, species that were dominant in one habitat tended to be fairly common across all habitats in the forest, suggesting that species–habitat preferences are relatively weak

Table 3 Dominant species based on stem abundance within the different habitats

Species	Family	Low depressions		Low flat		High gullies		Slopes		Ridge top		Full plot	
		N (ha)	Rank	N (ha)	Rank	N (ha)	Rank	N (ha)	Rank	N (ha)	Rank	N (ha)	Rank
<i>Phyllobotryon spathulatum</i> .(1)	Salicaceae	392.1	2	921	1	144	8	343	3	433	3	535	1
<i>Cola semecarpophylla</i> . (1)	Malvaceae	348.5	3	765.7	2	109	11	87.5	16	667	2	490	2
<i>Dichostemma glaucescens</i> (3)	Euphorbiaceae	271	4	443.6	3	213	5	248	5	423	4	345	3
<i>Cola praeacuta</i> (1)	Malvaceae	90.5	11	58.4	33	785	1	638	1	708	1	309	4
<i>Oubanguia alata</i> (3)	Lecythidaceae	469.6	1	355.2	4	90.5	16	60.4	23	126	11	298	5
<i>Cola</i> sp. Nov.2 (1)	Malvaceae	235.5	6	268	5	311	3	221	6	218	5	247	6
<i>Cola</i> sp. Nov.3 (1)	Malvaceae	239.8	5	200.9	6	51.4	23	37.7	31	92.5	18	165	7
<i>Diospyros preussii</i> (1)	Ebenaceae	138.3	8	123.6	11	169	7	110	12	209	6	147	8
<i>Angylocalyx oligophyllus</i> . (1)	Fabaceae	80.8	12	90.2	21	186	6	177	9	155	7	116	9
<i>Rinorea lepidobotrys</i> (1)	Violaceae	141.4	7	177.4	7	7.2	95	0.4	276	48.5	23	110	10
<i>Diospyros iturensis</i> (2)	Ebenaceae	42.6	34	89.3	22	102	13	214	7	136	9	95.9	11
<i>Rinorea subintegrifolia</i> (1)	Violaceae	60.3	25	88.2	23	98.1	15	196	8	109	14	94.1	12
<i>Rinorea gabunensis</i> (1)	Violaceae	3.1	173	3	177	342	2	383	2	125	12	94.1	13
<i>Cola cauliflora</i> (1)	Malvaceae	124.8	10	114.5	15	11.8	69	3.4	148	44.1	26	85.2	14
<i>Drypetes staudtii</i> (2)	Putrangivaceae	15.1	79	13.8	84	286	4	250	4	130	10	82	16
<i>Crotonogyne strigosa</i> (1)	Euphorbiaceae	65.4	21	143.2	9	13	66	50	26	37.2	30	77.7	19
<i>Tabernaemontana brachyantha</i> (3)	Apocynaceae	43.9	33	42.7	40	90.3	17	98.9	13	144	8	71.7	20
<i>Rinorea dentata</i> (1)	Violaceae	43.9	32	59.3	32	123	10	98.7	14	93.5	17	70	21
<i>Rinorea leiophylla</i> (1)	Violaceae	74	15	143.4	8	0.2	300	0	347	1.9	184	68.1	22
<i>Rinorea kamerunensis</i> (1)	Violaceae	61.1	24	130.3	10	4.9	118	12.1	79	36.6	31	68	23
<i>Protomegabaria stapfiana</i> (3)	Phyllanthaceae	137	9	31.9	50	78.9	20	16.3	70	24.7	47	67.3	24
<i>Strombosia</i> sp. (3)	Olacaceae	6.4	117	3.6	161	125	9	139	10	123	13	50.5	34

A species is considered dominant if it ranked among the top-10 in abundance in at least one of the five habitats. Species rankings in each habitat are listed. Densities in bold indicate significant species–habitat association according to torus-translation tests. The table is sorted by abundance rank in the entire plot, given in the last column. The life-form is given in parentheses (1 = treelets/small trees, 2 = understorey, 3 = lower canopy, 4 = upper canopy)

(i.e., growth and survival are only slightly reduced outside of the preferred habitat; Yamada et al. 2007) or that preferred habitats serve as a source of propagules for less optimal ‘sink’ habitats.

We found mixed support for our initial hypothesis that habitats with many associated species would be the most diverse. Low depressions had the most associated species, nearly twice as many as any other habitat, and also had the highest diversity per quadrat. At the other end of the spectrum, ridge tops, one of

the least diverse habitats, had the fewest species associations. However, the low flat habitat, which had the highest species richness per quadrat, had fewer species associations than the less diverse slope and high gully habitats.

The observed patterns of diversity are likely related to variation in soil moisture availability among habitats. The Korup FDP receives very high annual rainfall (>5000 mm/year), but experiences a severe, 3-month dry season. The well-drained soils of the

Table 4 Significant positive species–habitat associations of 272 species with ≥ 50 individuals in the Korup Forest Dynamics Plot based on torus-translation tests

Habitat	Low depressions	Low flat	High gullies	Slopes	Ridges
Low depressions	75				
Low flat	0	31			
High gullies	1	0	32		
Slopes	1	0	5	39	
Ridges	0	0	0	1	8

On the diagonal (in bold) are numbers of species associated with a given habitat. Below the diagonal are numbers of species associated with both the habitat given by the column heading and the habitat given by the row label for each cell

ridge top are most likely to be associated with water-stress in trees during the dry season, which may result in lower species diversity since drought-sensitive species typically have low survival in such habitats during the dry season (Comita and Engelbrecht 2009). In contrast, the lower habitats probably experience less water stress during the dry season, and may therefore support a higher diversity of trees. This is consistent with regional-scale trends of lower diversity of tropical trees in areas with lower moisture and longer seasonal droughts (Gentry 1988), and is thought to be because fewer species can physiologically tolerate stressful drought conditions. This idea is supported by our finding that very few species were associated with ridge tops, the driest habitat type. However, we also cannot rule out effects of soil nutrients, which vary spatially in the Korup plot (Chuyong, unpublished data) and can also influence tree species distributions and diversity (Newbery and Proctor 1984; Paoli et al. 2006; John et al. 2007).

Habitat associations and species coexistence in plant communities

We found that the majority of species in the Korup plot were significantly associated with at least one of the five topographic habitat types. Although partitioning at the coarse scale of habitats examined in this, and similar studies would clearly be insufficient for explaining the coexistence of the hundreds of species typically present in tropical tree communities, our results suggest that habitat partitioning plays a role in shaping species distributions and contributes to the maintenance of diversity in the Korup forest.

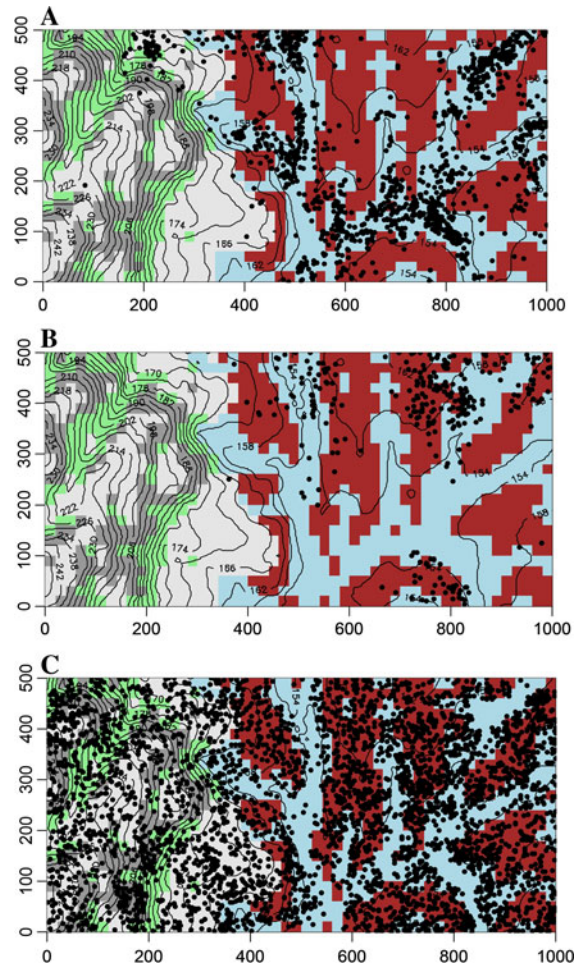


Fig. 3 Selected species distribution maps overlain on the habitat map of the Korup Forest Dynamics Plot. **a** *Beilschmiedia* sp. **b** *Lecomedoxa klaineana*. **c** *Hymenostegia afzelii*

Given the apparent variation in soil moisture of topographic habitats within the plot, the observed species–habitat associations provide indirect support for the idea of hydrologic niche partitioning (Silvertown et al. 1999; Webb and Peart 2000; Gibbons and Newbery 2003). Studies in a meadow community in Europe and a fynbos community in South Africa have revealed niche segregation along fine-scale hydrological gradients, due to a trade-off between species tolerance of aeration stress and drought stress (Silvertown et al. 1999; Araya et al. 2010). In tropical forests, experimental work has shown that tree species vary widely in their drought sensitivity (Engelbrecht and Kursar 2003), and that variation drives species distributions across both local and regional moisture gradients (Engelbrecht et al. 2007;

Baltzer et al. 2008). Similarly, species habitat associations within the Korup plot may be related to species distributions at regional scales. For example, the tree species *Oubanguia alata*, which was associated with low depressions in the plot, is mostly limited to wet forests of the Korup-Mount Cameroon area (Chuyong et al. 2004a, b). By contrast, both *Hymenostegia afzelii* (Fig. 3) and *Annickia affinis* are widely distributed in the plot and are also common in both wet and dry forests in the region (Aubréville 1970; Versteegh & Sosef 2007).

In Korup, strong habitat associations were shown predominantly by canopy and upper canopy species. This contrasts with findings from elsewhere. At Yansuni, Ecuador, significant habitat association was shown predominantly by understory species (Valencia et al. 2004), while in the dipterocarp forest at Sinharaja, Sri Lanka, all growth forms including most of the abundant species and most of the canopy dominants were habitat specialists (Gunatilleke et al. 2006).

Of the 172 species we tested, 63% showed significant positive habitat associations based on the conservative torus-translation test of association. This proportion is similar to the 64 and 79% obtained through similar torus-translation tests for Barro Colorado Island, Panama (Harms et al. (2001) and Sinharaja FDP, Sri Lanka (Gunatilleke et al. 2006), respectively. Philips et al. (2003) also reported an exceptionally high degree of habitat specialization (nearly 80%) for relatively well-sampled species in the Amazon, the highest record to date for tropical forest studies. Differences in the proportion of species showing significant habitat associations may be due either to variation in the strength of niche partitioning or to differences in the amount of topographic heterogeneity in the landscape captured within the plot area. Thus, it is difficult to make inferences about different processes acting in different forests based solely on proportion of associated species within a plot. Nonetheless, the high proportion of species associated with habitats in the Korup plot indicates that the lower diversity of African forests is unlikely to be due to lower topographic variation or less niche partitioning in these forests relative to forests in other tropical regions.

In conclusion, similar to patterns reported for tropical forests in Asia and the Americas, we found that topographic variation drives tree species distributions on local scales and plays a significant role in shaping

forest structure, species composition and diversity in this African forest plot. These results support the idea that niche partitioning contributes to the maintenance of diversity in species rich plant communities.

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