



Proceedings

Impact of Canopy Gap Ecology on the Diversity and Dynamics of Natural Regeneration in a Tropical Moist Semi-Deciduous Forest, Ghana *

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Abstract: The study evaluated species diversity, including regeneration and growth dynamics of different naturally regenerated tree species in gaps of different sizes in a tropical moist semi-deciduous forest, Ghana. Twenty-three randomly selected gaps categorized as small (<300 m²), medium (301-1000 m²), large (1001-2000 m²) and very large (>2000 m²) sizes were defined. Forty-one subsampling circular 1 m² plots at 2 m intervals were delineated within each gap. In total, 1468 individuals belonging to 85 species from 25 families and 65 genera were enumerated. Malvaceae (13 species) was the most diverse family, while Bignoniaceae, Clusiaceae, Lecythidaceae, Melastomataceae, Mimosaceae, Myristicaceae, Ochnaceae, Rutaceae, Santalaceae, and Urticaceae families were scarcely represented by only 1 species, respectively. Non-pioneer light demanding (NPLD) recorded the highest number of 38 species, followed by pioneers with 30 species, whereas shade-tolerant attained the lowest representatives of 17 species in regenerated species composition. All estimated diversity indices revealed improvements of species diversity in all gaps significantly. Though the small gaps showed no regeneration shift between pioneers and shade-tolerant, the other gap sizes did it at 62-82%. Medium and large gaps promoted natural regeneration at all distinguished diameter classes, however, pioneers and NPLD at 71-100 mm and pioneers at 51-70 mm were absent in small and very large gaps, respectively. Similarly, pioneers of height 0-20 cm in very large gaps were missing. Conclusively, species biodiversity differed significantly among gaps while regeneration dynamics differed significantly among gap sizes.

Keywords: diversity; dynamics; gaps; regeneration shift; tree species

1. Introduction

Canopy gaps are indispensable growing sites for the sustainable succession of different natural regeneration tree species. Generally, canopy gaps influence forest regeneration cycles by offering local resource-rich niches for understory tree species [1]. Differences among tree species in establishment, germination, development, survival, growth, and canopy recruitment within gap sites explain the crucial role of canopy gaps in shaping forest composition [1,2]. This very important role of canopy gaps had been well documented in numerous gap regeneration studies conducted in temperate [3] and tropical [4] forest ecosystems. Studies primarily focus on the effects of canopy gaps on species diversity [4,5], leaving only a few researching into the ecological positive effects of canopy gaps on natural regeneration dynamics [6]. Moreover, papers contributing to the widening knowledge on regeneration dynamics of naturally regenerated tree species with different shade tolerance mechanisms in gaps generally are quite common [6,7]. However, the role of gap size on ecological regeneration dynamics of natural regeneration has so far received

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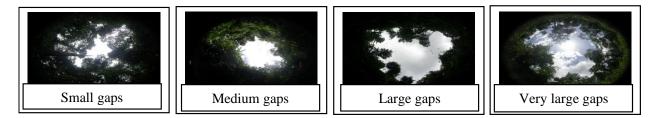


Figure 1. Canopy gaps of different sizes in a tropical moist semi-deciduous forest. Source: authors' own construct (2021).

2. Methods

The study was conducted in 84-year-old Bia Tano forest reserve, a tropical moist semi-deciduous forest in Ghana. The study area under prevailing tropical humid climate is located between latitude 6°52′–7°05′ N and longitude 2°30′–2°42′ W at 213 m–274 m a.s.l elevation with an area coverage of over 18, 000 ha. The mean annual air temperature and precipitation range between 26–29 °C and 1250–1750 mm, respectively. Schist and phyllites are the predominant underlying bedrocks with forest ochrosols as the most prevailing soil type according to Ghana soil classification system. The study area represents a hallmark of rich-species forest region in Ghana [4], a home of predominantly rich *Celtis*-*Triplochiton* (Cannabaceae-Malvaceae) association [4]. Twenty-three randomly selected gaps categorized as small (n = 2; ≤300 m²), medium (n = 7; 301–1000 m²), large (n = 11; 1001–2000 m²) and very large (n = 3; >2000 m²) sizes were defined (see Table 1). Forty-one subsampling circular 1 m² plots at 2 m intervals were delineated within each gap. Every naturally regenerated tree species of height 0–350 cm and diameter 0–100 mm within subsampling plots were identified by species and recorded.

Paleontological Statistics software (3.24) [8] was used to estimate various assessed biodiversity indices. ANOVA and *post hoc* Tukey's HSD test were carried out to show significance of means at p < 0.05 level using STATISTICA (13.4.0.14). Simpson's index (1-D) measured species dominance, Shannon diversity index (H) measured species diversity, Berger-Parker Index (B-P) measured most abundant species; Pielou's Evenness (J) measured species evenness, Menhinick's index (Mk); Fisher's alpha (α) and Chao1 diversity index (C-1) measured species richness from different perspectives (Figure 2). Formulas for different analyzed biodiversity indices based on adopted Equations (1), (2), (4)–(7) [9] and (3) [10] are displayed in Figure 2.

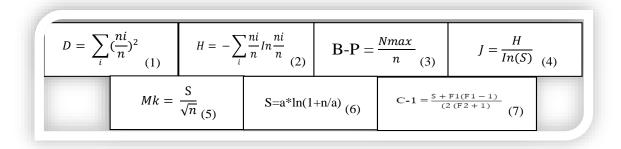


Figure 2. Plate of 7 assessed biodiversity indices (where, S is the number of taxa, n is the total number of individuals, D is Dominance, n*i* is the number of individuals of taxon ith, in is the logarithm, a is Fisher's alpha, Nmax is the total number of individuals of the most abundant species, F1 is the number of singly regenerated species and F2 is the number of doubleton regenerated species).

Table 1. Presenting Total individuals (N), Taxa (S), Simpson's index (1-D), Shannon diversity index (H), Berger-Parker
dominance (B-P), Menhinick richness index (Mk), Pielous Evenness (J), Fisher alpha (a) and Chao-1 (C-1) biodiversity
indices to express species diversity of naturally regenerated tree species in gaps of different sizes at tropical moist semi-
deciduous forest in Ghana.

	Estimated Biodiversity Indices									
Gap ID (Size/Area m ²) -		Ν	S	1-D	Н	B-P	J	Mk	α	C-1
Gap1(Large/1896)		123	29	0.87	2.60	0.29	0.78	2.49	11.14	29.41
Gap2(Medium/782)		90	16	0.56	1.49	0.65	0.54	1.62	5.32	24.78
Gap3(Medium/782)		57	11	0.65	1.56	0.55	0.65	1.46	4.06	17.72
Gap4(Large/1252)		35	13	0.85	2.18	0.26	0.87	2.08	6.83	28.08
Gap5(Large/1098)		46	17	0.87	2.39	0.25	0.86	2.36	8.78	34.62
Gap6(Very large/2557)		102	14	0.77	1.86	0.36	0.72	1.32	4.11	20.57
Gap7(Large/1244)		48	15	0.85	2.27	0.28	0.84	2.12	7.21	22.32
Gap8(Large/1457)		12	9	0.84	1.96	0.25	0.94	2.31	12.50	14.03
Gap9(Medium/692)		25	13	0.84	2.18	0.29	0.88	2.40	9.34	25.67
Gap10(Large/1249)		52	11	0.81	1.95	0.31	0.81	1.48	4.07	17.67
Gap11(Large/1425)		73	15	0.79	2.03	0.37	0.76	1.64	5.18	22.67
Gap12(Large/1420)		37	9	0.83	1.94	0.27	0.89	1.48	3.79	12.00
Gap13(Large/1647)		33	9	0.73	1.63	0.39	0.77	1.45	3.62	13.50
Gap14(Large/1058)		39	15	0.86	2.29	0.25	0.87	2.24	7.92	23.31
Gap15(Very large/2728)		20	9	0.83	1.95	0.27	0.90	1.94	5.84	14.42
Gap16(Large/1664)		50	19	0.88	2.48	0.23	0.86	2.55	10.17	34.70
Gap17(Very large/2005)		57	19	0.90	2.60	0.18	0.89	2.43	9.39	27.78
Gap18(Small/296)		44	18	0.91	2.62	0.17	0.91	2.61	10.60	24.67
Gap19(Small/287)		170	19	0.79	2.05	0.37	0.70	1.46	5.48	23.38
Gap20(Medium/607)		108	29	0.91	2.81	0.18	0.84	2.76	12.75	42.05
Gap21(Medium/884)		88	15	0.77	1.89	0.37	0.71	1.49	4.72	21.17
Gap22(Medium/971)		100	20	0.71	1.98	0.51	0.66	2.00	7.52	37.00
Gap23(Medium/837)		59	17	0.80	2.13	0.37	0.76	2.08	7.27	31.94
	df			22	22	22	22	22	22	22
	F-ratio			5.94	7.52	4.21	6.04	15.37	7.72	2.15
	<i>p</i> -value			***	***	***	***	***	***	*

Total number (n) presents individual species. Taxa (S) is the number of presented species. p < 0.001 (***) significance level, p < 0.05 (*) significance level.

3. Results and Discussion

3.1. Impact of Gap Ecology on Species Composition and Diversity

In total, 1468 individuals belonging to 85 species from 25 families and 65 genera were enumerated. Malvaceae (13 species) was the most diverse family, while Bignoniaceae, Clusiaceae, Lecythidaceae, Melastomataceae, Mimosaceae, Myristicaceae, Ochnaceae, Rutaceae, Santalaceae, and Urticaceae families were scarcely represented by only 1 species, respectively. This result firstly demonstrates the species-rich diversified nature of tropical forests and further validates the importance contribution of tropical forests towards the maintenance of global tree biodiversity [11]. Secondly, this observation clearly depicts that gaps remain the conducive habitats for rich species composition of natural regeneration [1,7]. Non-pioneer light demanding (NPLD) recorded the highest number of 38 species, followed by pioneers with 30 species, whereas shade-tolerant attained the lowest representatives of 17 species in regenerated species composition. This finding supports the long-standing notion that light-tolerant species prefer open space environments for optimum regeneration [5] while those environments normally impede regeneration density, relative

density = 97, 774 trees/ha, 6.61%) and *Daniella ogea* (85, 678 trees/ha, 5.79%), NPLD- *Mansonia altissima* (75, 598 trees/ha, 5.11%), and *Blighia sapida* (70, 558 trees/ha, 4.77%), and shade-tolerant- *Nesogordonia papaverifer* (340, 2711 trees/ha, 23.16%), *Celtis mildbraedii* (188, 1499 trees/ha, 12.81%) and *Baphia nitida* (185, 1475 trees/ha, 12.60%) were the predominant tree species found almost in all gaps. In another tropical moist semi-deciduous forest in Ghana, *Albizia zygia* (0.50–52.96%), *Ceiba pentandra* (0.67–75.67%), *Celtis mildbraedii* (0.74–22.16%) and *Turreanthus africanus* (0.67–33.16%) among 37 encountered natural regeneration species were the four predominant species that significantly influenced high seedling densities during 0.5, 10 and 33 months in gaps following selective logging [6].

Furthermore, all estimated biodiversity indices revealed improvements of species diversity in all gaps significantly (Table 1). Nonetheless, biodiversity did not vary greatly from gap to gap (287–2728 m²). Thus, our finding is consistent with Schnitzer and Carson [5] that gaps play a similar role in maintaining tree species diversity in many tropical forests. By contrast, disprove gap size as a significant variable on naturally regenerated tree species composition and diversity according to Hammond et al. [3]. Generally, this observation solidly corroborates a view that gap-based silviculture plays a vital role in forests by promoting ecological species assemblage mechanisms that enable species coexistence which significantly relates to the positive impacts of species diversity on forest ecosystem's stability and functions [3].

3.2. Impact of Gap Ecology on Regeneration Dynamics of Species with Different Shade Tolerance Mechanisms

Our analysis investigating the ecological regeneration shift of pioneers to shade-tolerant species in gaps of different sizes. There, from small to medium, large and very large gaps, the abundance proportions among Pioneers (%): Non-pioneer light demanding species (%): Shade-tolerant tree species (%) roughly varied as: 39:18:43, 13:19:68, 19:20:61 and 10:24:66, respectively. Trends showed that the rate of sequential replacement of pioneers by shade-tolerant species within very large gaps (+4.94) was highly rapid compared to medium gaps (+5.86) and large gaps (+13.12) whereas no ecological species replacement was found within small gaps (+33.70). Besides, a stronger replacement relationship between concerned species within very large gaps (82%) and medium gaps (72%) areas were observed, while a less weak ecological association was detected within large gaps (62%). Still, our results substantiate the long-established knowledge about regeneration shift between pioneers and shade-tolerant species in gaps under tropical forests [4,6,7] due to species biological growth characters and physiological light tolerance ability; in addition, the colossal size of very large (average size 2430 m²) and large (1401 m²) gaps, and higher light levels in medium gaps made this natural process more rapid and effective.

Again, from Figure 3, no significant difference (p > 0.05) was found between species with different shade mechanisms in small and very large gaps. However, there were significant differences among tree species at p < 0.001 level in medium and large gaps. But then, no significant difference (p > 0.05) was observed between the two light-tolerant species. In addition, higher proportions of shade-tolerant species in all gap sizes were observed. Hence, the imbalance proportions of 2:1:2, 1:2:5, 1:1:3 and 1:2:6 were observed among pioneers: NPLD: shade-tolerant species in small, medium, large, and very large gaps, respectively. Gaps maintaining dominated shade-tolerant species originated from either ecological regeneration replacement or were present as advance regeneration prior to gap creation. Therefore, our findings are consistent with those of [4] and [5] specifically, that gaps play a neutral role in maintaining tree species with different lifelong light demanding histories, however, abundance proportions for only shade-tolerant species are usually high. Small gaps showing similar proportions between pioneers (1459 trees/ha) and shade-tolerant (1573 trees/ha) species suggest that gaps of relatively smaller sizes (<300 m²) are essential growing areas for preserving balanced proportions of natural regeneration of different suites of tree species.

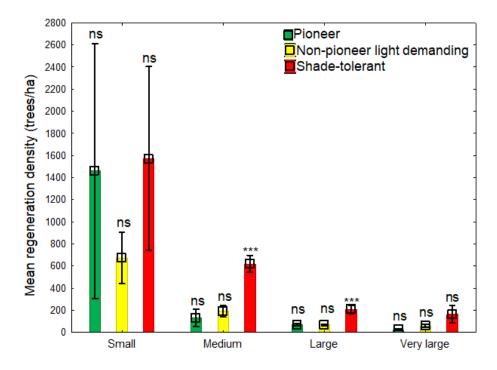


Figure 3. Multiple comparison of different naturally regenerated tree species in small (n = 2); medium (n = 7); large (n = 11) and very large (n = 3) gaps.

3.3. Impact of Gap Ecology on Growth Dynamics of Different Naturally Regenerated Tree Species

From Figure 4 (left plate), it was observed that medium and large gaps promoted natural regeneration at all diameter classes (0–10, 11–30, 31–50, 51–70, and 71–100 mm), however, pioneers and NPLD at 71–100 mm and pioneers at 51–70 mm were absent in small gaps, and pioneers at 51–70 mm were again absent within very large gaps. Similarly, pioneers at 0–20 cm in very large gaps were missing in regeneration height growth dynamics in Figure 4 (right plate).

The missing light-tolerant representative species at the two sapling classes (51–70 and 71–100 mm) of diameter growth dynamics justifies notion that small gaps are unyielding grounds for optimum growth of light dependent tree species. The full appearance of all height classes (0–20, 21–50, 51–100, 101–200, and 201–350 cm) for categorized tree species groups in almost all gap sizes under height growth dynamics in comparison to absent diameter classes of diameter growth dynamics strongly attests to the finding in several studies that in tree growth under prevailing light environments, height growth is often maintained over diameter growth, e.g., [12]. But then again, our result disputes the assumption that differences in the light environments of trees growing under various levels of canopy cover would be expressed sooner in diameter growth than they would in height growth [13,14].

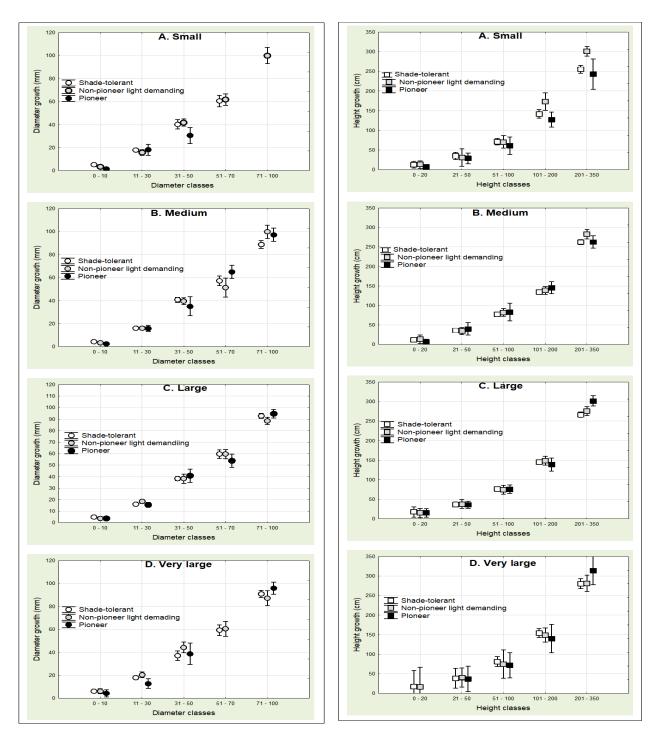


Figure 4. Multiple comparison of diameter (left) and height (right) growths of different naturally regenerated tree species in small (n = 2); medium (n = 7); large (n = 11) and very large (n = 3) gaps.

4. Conclusions

Conclusively, the present study demonstrates some beneficial effects of canopy gap ecology on the diversity and dynamics of natural regeneration in a moist tropical forest and adds a valuable contribution to tropical forestry, especially in Ghana and tropical African regions. Species diversity differed significantly among gaps but not among gap sizes. Expansive size of canopy gaps and higher light levels in gaps were the explanatory factors for the rapid ecological regeneration shift of pioneers to shade-tolerant species in medium to very large gaps at 62–82%. Higher proportions of shade-tolerant species (43–68%) in all

gap sizes were observed. Gap size was a significant variable that affected species dynamics of natural regeneration in gaps. Small gaps were found to be unyielding grounds for light dependent species in sapling classes (51- 70 and 71–100 mm) of diameter growth dynamics. Gaps as growing sites promoted the full appearance of all height classes (0–20, 21–50, 51–100, 101 -200, and 201–350 cm) for all categorized gap regeneration tree species.

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