SPATIAL PATTERNS AND DEMOGRAPHICS OF *STREBLUS MACROPHYLLUS* TREES IN A TROPICAL EVERGREEN FOREST, VIETNAM

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NGUYEN H, WIEGAND K & GETZIN S. 2014. Spatial patterns and demographics of *Streblus macrophyllus* trees in a tropical evergreen forest, Vietnam. *Streblus macrophyllus* is a shade-tolerant and subcanopy tree species common to tropical evergreen forests in northern Vietnam. However, its ecology is poorly known. We used spatial point pattern analysis to describe the spatial arrangement of tree individuals within a forest community dominated by *S. macrophyllus*. All individual trees with diameter at breast height larger than 2.5 cm in a 1-ha plot were mapped and measured. The overall pattern of this species was a regular distribution at scales up to 2 m. Its juveniles and subadults were strongly aggregated, but adult trees were regular at scales of up to 3 m, implying evidence of density dependent thinning. The spatial pattern of *S. macrophyllus* strongly affected the patterning of the whole plot. In *S. macrophyllus*, juveniles and subadults were similarly distributed relative to adults and showed additional clumping independent of the adults. The overall interspecific association between adults of other species and *S. macrophyllus* at different life-history stages also showed independence. We conclude that *S. macrophyllus* is a predominant competitor within the community and it follows a gap-phase regeneration mode.

Keywords: Point pattern analysis, forest community, gap-phase regeneration mode, intra- and interspecific associations

INTRODUCTION

Streblus macrophyllus (Moraceae) is a common tree species in the tropical rainforests of northern Vietnam. *Streblus macrophyllus* has synonyms with some other scientific tree names such as *Taxotrophis balansae*, *Dimerocarpus balansae* and *Dimerocarpus brenieri* (Ho 1999, eFlohras 2010). The species is a subcanopy tree with height of up to 20 m. In Cucphuong National Park, Vietnam, *S. macrophyllus* is dominant in the humid valleys of the limestone mountains. However, the ecology of this species is poorly known.

Trees are too long-lived for feasible study of the dynamics of cohorts from sapling to adult stage. Thus, it is easier to use the size-structured approach to investigate forest population dynamics (Turner 2004). Spatial patterns of cohorts can reflect contemporary or historical processes affecting the population. Therefore,

ecological processes in plant ecology such as dispersal, competition, survival and growth can be elucidated with the application of spatial statistics (Penttinen et al. 1992, Stoyan & Penttinen 2000).

We used recent techniques of spatial point pattern analysis to detect spatial patterns of trees and test null hypotheses that were related to factors controlling spatial arrangements of trees (Diggle 2003, Wiegand & Moloney 2004). The variation of tree density at different spatial scales can be observed and may be determined by separate causes. It is commonly observed in a given species that spatial patterns shift from clumping to regularity with increasing tree size and this may be caused by density-dependent thinning explained by intraspecific competition for resources (John et al. 2002, Stoll & Bergius

2005, Freckleton & Lewis 2006) or only by high overall mortality (Wiegand et al. 2000). Moreover, negative associations between adult and early life-history stages, e.g. juvenile and subadult, provide useful information to describe regeneration processes of the species such as colonisation following dispersal or density-dependent mortality (Grau 2000).

The goals of this study were to characterise and interpret spatial patterns of *S. macrophyllus.* Understanding the spatial structure and association of trees in different life-history stages may elucidate the processes controlling the dominance of this species in this forest. We hypothesised that intraspecific competition and clustered regeneration would be the dominant processes that facilitated species coexistence in general and abundance of the *S. macrophyllus* species in particular.

MATERIALS AND METHODS

Study site and data collection

The study was conducted at Cucphuong National Park, northern Vietnam. The national park was established in 1962 and covers an area of 22,200 ha (Figure 1). It is surrounded by limestone mountains with mean maximum height of 300–400 m and is covered by tropical evergreen rainforest. In the core zone, mean annual temperature is 20.6 °C, but mean temperature in winter is only 9 °C. In the buffer zone, mean annual temperature is about 2° higher. Annual mean humidity is 85% and the average annual rainfall is 2138 mm year^{-1}. The study site, with latitude of 20° 17' N, longitude of 105° 39 ' E, and elevation of 266 m above sea level was specifically selected to investigate
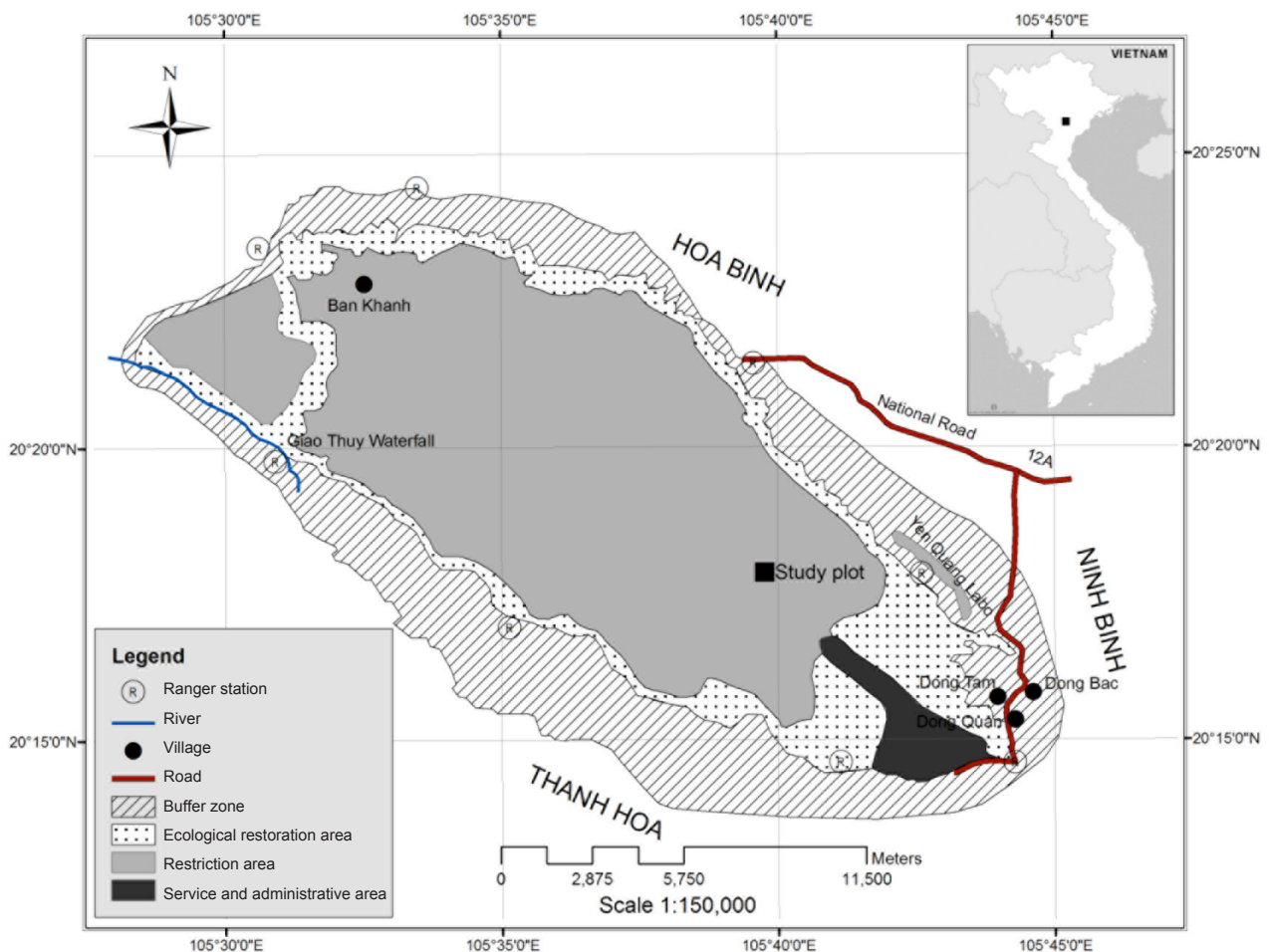

Figure 1   Map of Cucphuong National Park, Vietnam

the spatial distribution of *S. macrophyllus*. Its adult trees may reach about 15 m in height and the species is found in combination with other species of Clusiaceae, Ulmaceae and Annonaceae (FIPI 1996).

A 1-ha plot was created in an evergreen forest stand dominated by *S. macrophyllus* trees in the core zone. The plot was subdivided into a grid of 100 subplots of 10 m × 10 m. All woody plants with diameter at breast height (dbh, at 1.3 m above ground) ≥ 2.5 cm were identified to species, stem-mapped to their relative geographical coordinates (x, y) and their dbh recorded to the nearest 0.5 cm using a laser distance measurement device, compass and diameter tape. All individual trees were allocated into one of the three basic life-history stages, namely, juvenile (dbh < 5 cm), subadult (5 cm ≤ dbh < 10 cm) or adult (dbh ≥ 10 cm).

Statistical analysis

The pair-correlation function and Ripley's K-function are based on the distribution of distances of pairs of points (e.g. x, y coordinates of trees). Both functions are standard methods for analysing mapped point patterns in ecology (Condit et al. 2000, Wiegand et al. 2007) particularly in forest ecology used to quantify spatial patterns of tree species. Ripley's K-function is defined as the expected number of points within distance r of an arbitrary point divided by the intensity λ where λ is the intensity of the pattern in the study area (Ripley 1976). We used the L-function (Besag 1977), a transformation of Ripley's K-function, $L(r) = (K(r)/\pi)^{0.5} - r$. The pair-correlation function is the derivative of the K-function (Stoyan & Stoyan 1994, Illian et al. 2008), $g(r) = K'(r)/(2\pi r)$. Particularly, it is non-cumulative and is defined as the expected density of points at a distance radius r from a randomly chosen point.

Based on the tree–tree distances, the univariate pair-correlation function $g_{11}(r)$ describes the spatial distribution of trees at a given radius r using a standardised density. Consequently, $g_{11}(r)$ = 1 under complete spatial randomness, $g_{11}(r) > 1$ indicates aggregation and $g_{11}(r) < 1$ indicates regularity at distance r within trees of the pattern. The bivariate pair-correlation function $g_{12}(r)$ was extended to describe spatial association between two types of points, for example two tree species. Function $g_{12}(r)$ was defined as the expected density of points of type 2 at distance r from an arbitrary point of type 1. Similarly to the univariate version, $g_{12}(r)$ = 1 indicates independence, $g_{12}(r) < 1$ indicates repulsion and $g_{12}(r) > 1$ indicates attraction between two tree species at distance r.

The null models of random distribution, spatial independence and random labelling of trees were tested for null hypotheses related to ecological questions (Goreaud & Pelissier 2003, Wiegand & Moloney 2004, Illian et al. 2008). Significant departure from the null models was evaluated using 999 Monte Carlo simulations and approximately 99% confidence envelopes were built by 5th lowest and 5th highest values of these simulations. All point pattern analyses were performed using the grid-based software Programita (Wiegand & Moloney 2004).

We used the null model complete spatial randomness to investigate whether the tree distribution is spatially restricted by environmental heterogeneity based on the spatial distribution of all adult trees. The distribution of all adult trees at scales beyond direct tree–tree interaction should indicate environmental suitability for colonisation and growth as well as allow capturing of environmental effects common to all species within the stand (Getzin et al. 2008).

Using the L-function and pair-correlation function allowed us to observe the change in tree density from both cumulative and non-cumulative perspectives at various scales. In plant communities, clumping at small scales generally indicated plant–plant interactions, while at large scales it showed the effects of environmental heterogeneity (Wiegand et al. 2007). Under heterogeneous site conditions, the pair-correlation function fluctuated above and below the upper simulation envelope, while the L-function indicated heterogeneity by a strong increase at large scales caused by highly cumulative density at those scales (Wiegand & Moloney 2004). However, in our plot, contrasting the pattern of all adult trees (dbh ≥ 10 cm) to the complete spatial randomness null model showed that there was no significant departure from the null model of complete spatial randomness at large scales (Figure 4). Consequently, the hypothesis of environmental homogeneity could not be rejected and we applied the homogeneous g-function for the following spatial pattern analyses.

Analysis 1: Overall univariate patterns

The spatial distributions in size classes of all species, *S. macophyllus* and the remaining species were analysed and compared using the univariate pair-correlation function. We tested the null hypothesis of a random univariate spatial distribution of trees in each life-history stage (juvenile, subadult and adult). If a random distribution of a life stage over the entire plot was confirmed, this meant that no strong interaction occurred between the trees within this size class. Comparisons of the spatial distributions of the whole plot with and without considering *S. macophyllus* allowed us to interpret the abundance effect of the species. Overall, we expected that spatial patterns of adult trees were more regular than those of earlier life-history stages, with the increase of tree size being evidence of self-thinning.

Analysis 2: Dispersion patterns of S. macrophyllus

Using random labelling null model, we investigated whether or not a random structure of two labels occurred within the joined patterns. The g-functions are invariant under random thinning of the joined patterns 1 and 2 (Wiegand & Moloney 2004). Thus, we expected $g_{12}(r) = g_{21}(r) = g_{11}(r) = g_{22}(r)$. In our case-control design, control was pattern 1 (adult trees) and case was pattern 2 (one of the early life-history stages). If the test statistic $g_{12}(r) - g_{11}(r) \approx 0$, it meant that individuals of pattern 2 around individuals of pattern 1 at scale r were distributed in the same way as individuals of pattern 1 around themselves. Therefore, this would imply that both life-history stages exploit the natural resources at that scale in a similar way (Getzin et al. 2008). In addition, if the test statistic $g_{21}(r) - g_{22}(r) << 0$, there were more individuals of cases than controls in proximity of case individuals, therefore we would expect an additional clustered pattern of cases independent of the controls (i.e. adult trees).

We used a case-control design to estimate mutual effects between early life-history stages (cases) and the adult stage (control) of *S. macrophyllus*. Under the null hypothesis of random labelling, if the cases did not show any significant difference from the controls then they were a random subset of the joint pattern of both cases and controls. We expected a clustering of the cases reflecting regeneration mechanisms via independent clumping relative to the controls and, if clustering decreased with increasing size class, density-dependent thinning with increasing size class. For example, there may be an additional clustering of juveniles independently from adult trees that may be caused by canopy gaps with conditions suitable for survival and growth of juveniles.

Analysis 3: Hetero-specific association of S. macrophyllus

To describe the spatial association between adult trees of the remaining species (pattern 1) and the different life-history stages of *S. macrophyllus* (pattern 2), we used the bivariate pair-correlation function. The null hypothesis of spatial independence was used to test for non-randomness in spatial arrangement between two patterns based on the assumption that the two patterns were generated by two independent processes. We kept the first pattern unchanged and then randomly shifted the second pattern relative to pattern 1 (Wiegand & Moloney 2004). We hypothesised that *S. macrophyllus* had gap-phase regeneration mode, i.e. the early stages of *S. macrophyllus* were clustered independently of con-specific adults and simultaneously, they were repulsed by hetero-specific adults. Thus, we expected a negative association of pattern 2 (a life-history stage of *S. macrophyllus*) relative to pattern 1 (adult trees of the remaining species).

RESULTS

Species composition and characteristics

A total of 705 individuals with dbh ≥ 2.5 cm were enumerated in the 1-ha study plot. Forty species were identified and belonged to 21 families while one species was unknown (Lauraceae). *Streblus macrophyllus* (Moraceae) was most abundant with 483 individuals ha^{-1} (Table 1, Figure 2), approximately 68.5% of the total. The second and third most abundant species were *Hydnocarpus kurzii* (Flacourtiaceae) and *Caryodaphnopsis tonkinensis* (Lauraceae) with 33 and 29 trees ha^{-1} respectively. Consequently, all species other than *S. macrophyllus* were grouped into 'remaining species' in our spatial pattern analyses.

Table 1   Stand structure and species composition

Species	Stem density (ha⁻¹)	Basal area (m² ha⁻¹)	Max dbh (cm)	Median NN (m)
Streblus macrophyllus	483	9.38	50.6	1.59
Hydnocarpus kurzii	33	1.20	42.9	4.03
Caryodaphnopsis tonkinensis	29	15.21	133.7	5.68
Others (38 species)	160	10.60	155.1	1.99
All	705	36.39	155.1	1.38

Dbh = diameter at breast height, NN = nearest neighbour distance
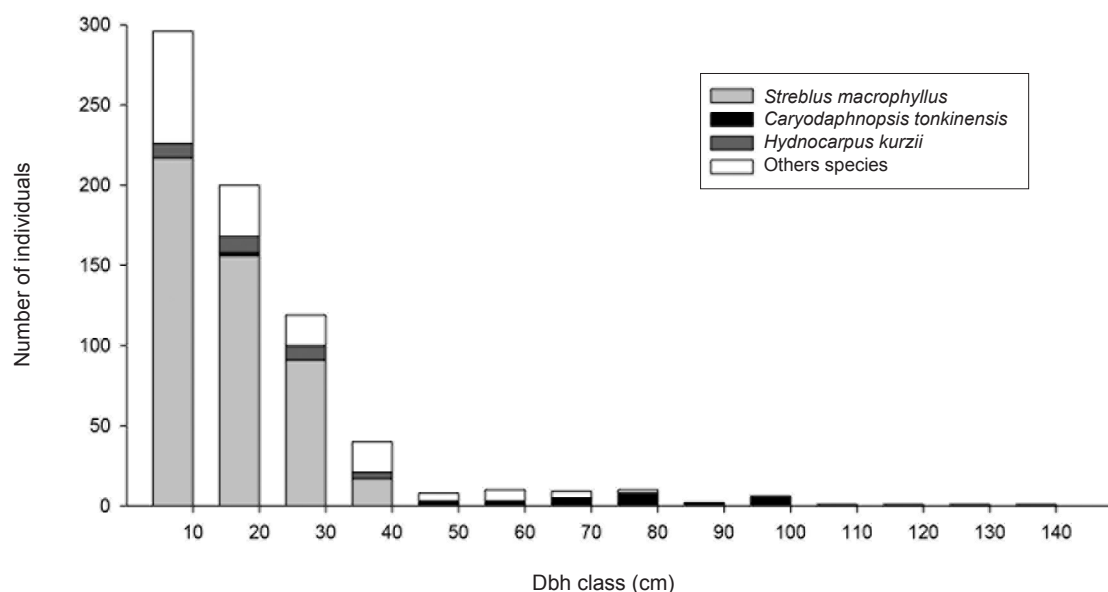
Figure 2   Diameter at breast height (dbh) class distribution of *Streblus macrophyllus* and the remaining species in the study plot

The diameter distribution of *S. macrophyllus* also showed a modal curve with a reverse J-shape, suggesting that the population regenerated continuously (Figure 2). This is a typical feature for uneven-aged and multi-species stands in tropical rainforests (Richards 1996). Considering all remaining species together, the size distribution was a reversed J-shape with a 'long foot'. *Hydnocarpus kurzii* and *C. tonkinensis* contributed small numbers of trees in large tree size classes (Table 1). Even though *S. macrophyllus* was the most abundant species, *C. tonkinensis* dominated in basal area with coverage of 15.21 m² ha⁻¹ due to its large size (Table 1, Figure 3).

Spatial analysis

Overall univariate patterns

Spatial pattern of all adult tress (dbh ≥ 10 cm) was regular at scales of up to 3 m, which was evidence of strong competition within adult trees (Figure 4b). Spatial patterns of all species clearly shifted from aggregation towards regularity with increasing tree size (Figures 5b–d). Combined wit regular patterns of all trees at scales up to 2 m (Figure 5a), these patterns showed that strong inter-tree competition occurred in this forest stand.

Similarly, spatial pattern of *S. macrophyllus* showed regular distribution at scales up to 1.5 m (Figure 5e). Its juveniles and subadults were clustered but intensities and scales decreased with increasing tree size (Figure 5f, g), while adult trees were regular up to scales of 3 m (Figure 5h).

Dispersion patterns of S. macrophyllus

The test statistic $g_{12}(r) - g_{11}(r)$ did not differ significantly from zero when comparing distributions of adults relative to those of
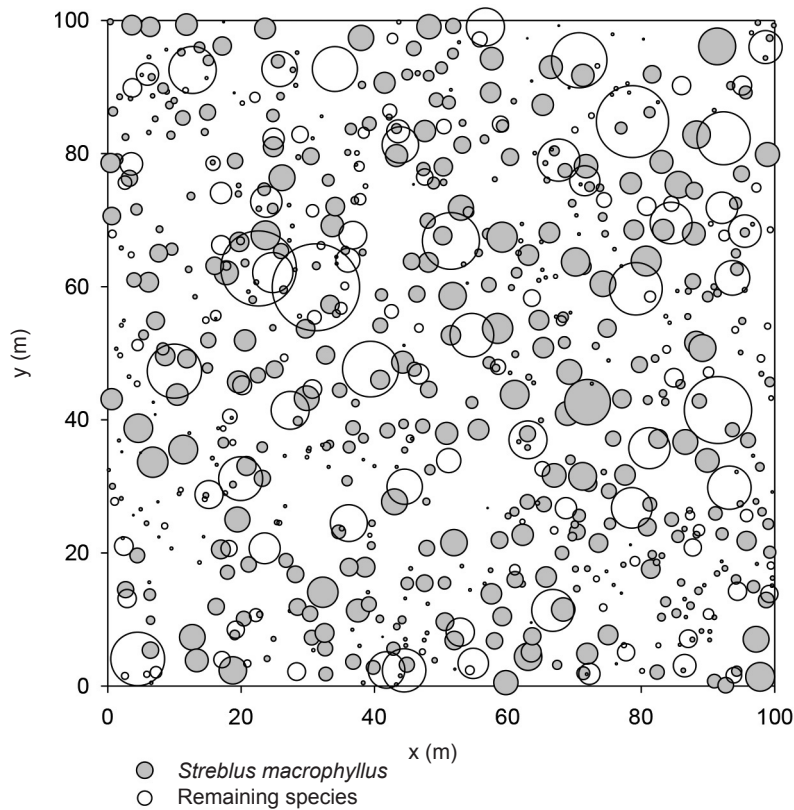
Figure 3　Distribution map of all tree individuals: *Streblus macrophyllus* (grey circles) and the remaining species (open circles); the size of symbols is proportional to the dbh of the individuals with an actual size of 2.5 to 155 cm; dbh = dimeter at breast height
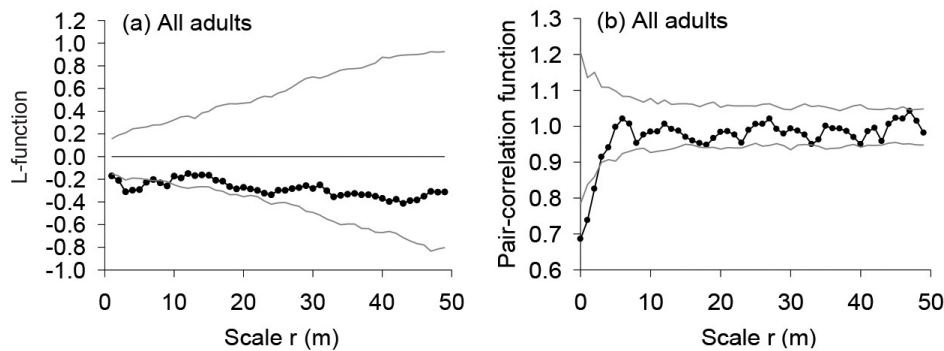

Figure 4　The univariate L-function and pair-correlation function show spatial distributions of all adult trees (dbh ≥ 10 cm) at different scales; the observed patterns (dark line) lying beyond the confidence envelopes (grey lines) indicate significant departure from the null model of complete spatial randomness

juveniles and subadults (Figures 6a and b). This meant that the distribution of juveniles or subadults surrounding adults did not differ from the distribution of adults relative to adults. Therefore, these results indicated that distribution of *S. macrophyllus* individuals in juveniles and subadults around adults $g_{12}(r)$ followed the same overall pattern of adult trees relative to themselves $g_{11}(r)$. These distribution patterns suggested that the succession process of *S. macrophyllus* was spatially homogeneous throughout all life-history stages.

Test statistic $g_{21}(r) - g_{22}(r)$ showed significant differences from the null model of random labelling for juveniles and subadults (Figure 6c and d). Results showed that there were strong
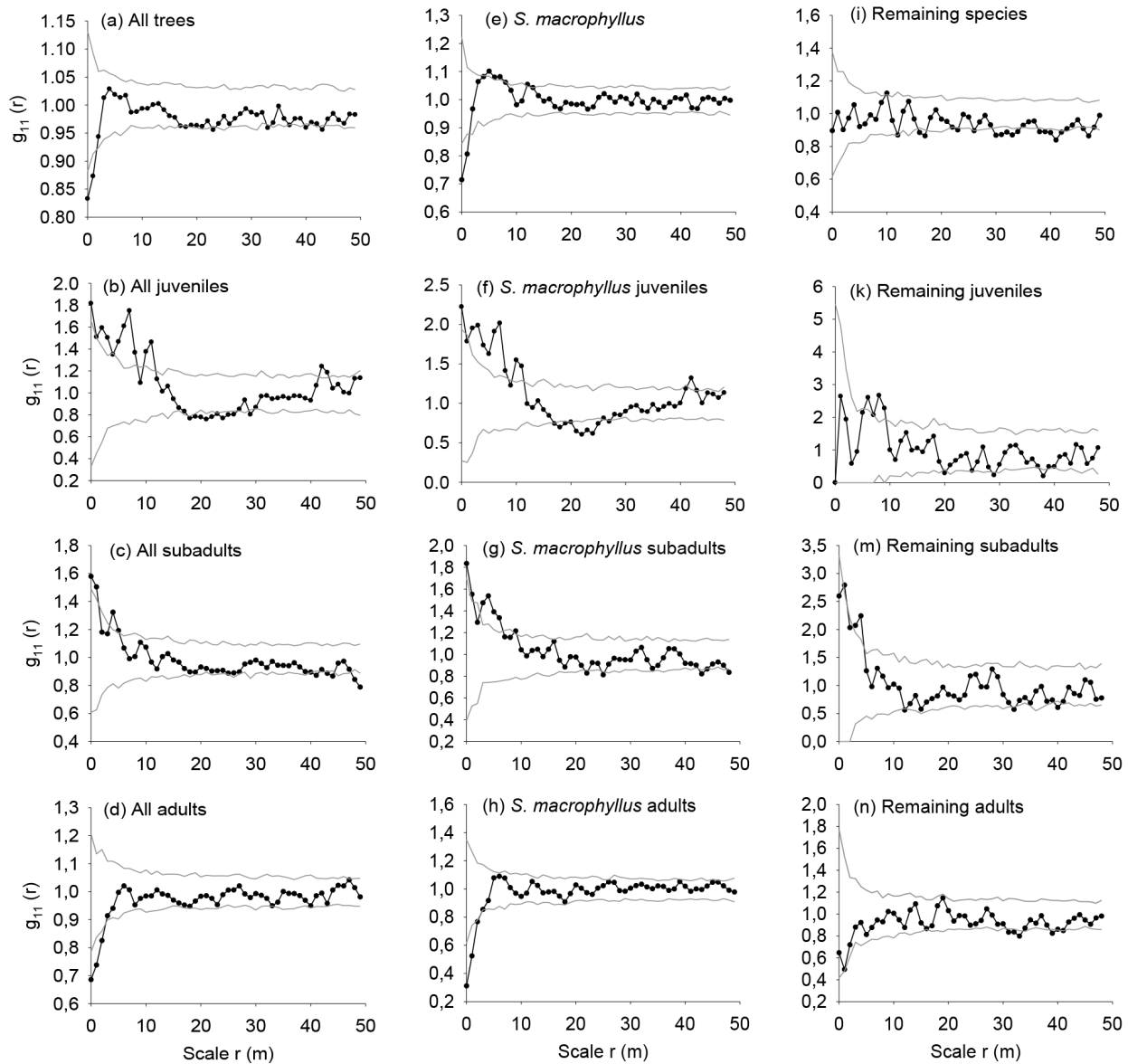
Figure 5   Spatial distributions of all individuals and life-history stages of all species (5a–d), *Streblus macrophyllus* (5e–h) and remaining species (5i–n) are shown by the univariate pair-correlation function; the observed patterns (dark line) lying beyond the confidence envelopes (grey lines) indicate significant departures from the null model of complete spatial randomness

additional clumpings which were independent of adult trees at fine scales up to 11 m in the case of juveniles and 6 m in subadults. At scales 18–24 m, $g_{21}(r) - g_{22}(r) > 0$ meant that there were significantly less juveniles surrounding adults compared with the adult–adult relationship (Figure 6c). Intensity and range of aggregation decreased with increasing tree size (life-history stages). This is evidence of increased intraspecific competition between young conspecific trees which were growing away from adults. These results were also compatible with inferences about the aggregation of young trees (Figures 5f

and g) and they showed that these aggregations were independent of the pattern of adult trees.

Heterospecific association of S. macrophyllus

Streblus macrophyllus juveniles were marginally repulsed by heterospecific adults at a scale of 7 m (Figure 7a). Subadults and adults of *S. macrophyllus* were independent of the heterospecific adults (Figures 7b and c). Thus, we did not find very strong evidence of interspecific repulsion to support the assumption of a gap-phase regeneration mode of *S. macrophyllus*.
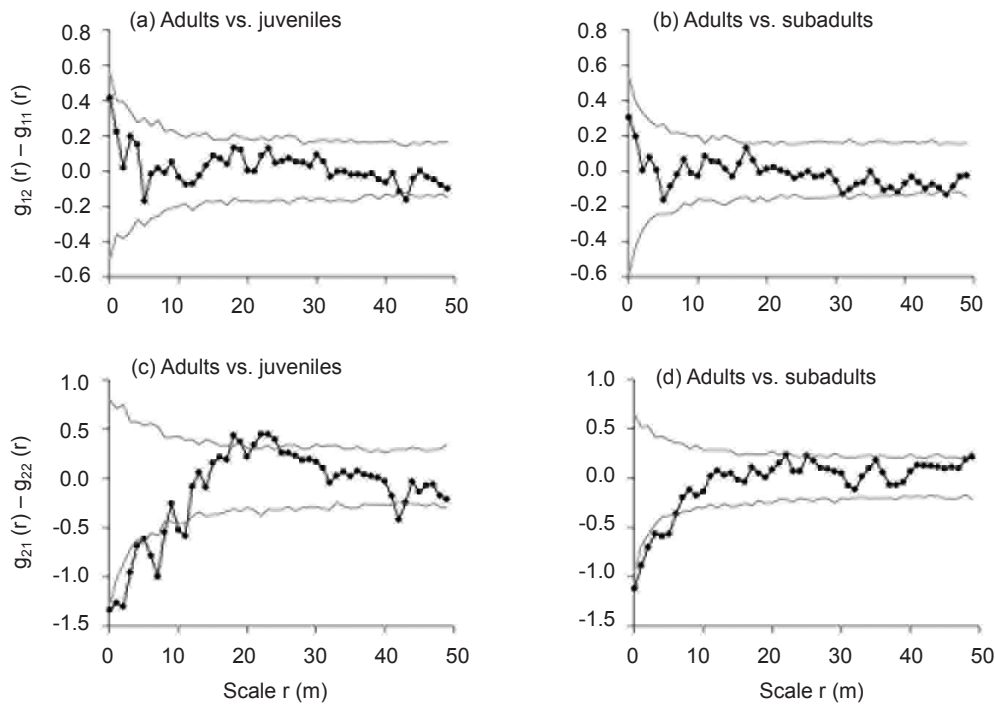
Figure 6　　Test statistics $g_{12}(r) - g_{11}(r)$ and $g_{21}(r) - g_{22}(r)$ showing spatial associations between adult stage and two early stages in the life-history of *Streblus macrophyllus*; the observed patterns (dark line) lying beyond the confidence envelopes (grey lines) indicate significant departures from the null model of random labelling
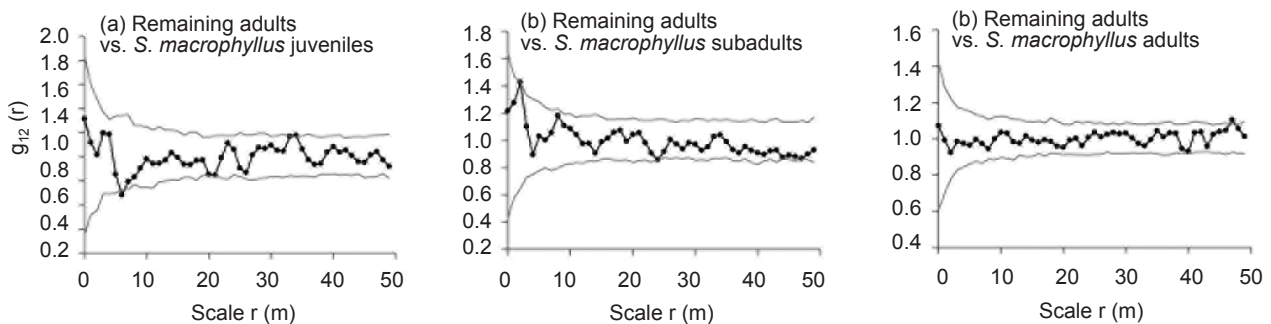

Figure 7　　The bivariate pair-correlation functions show association patterns between adults of remaining species and different life stages of *Streblus macrophyllus*; the observed patterns (dark line) lying beyond the confidence envelopes (grey lines) indicate significant departures from the null model of spatial independence

DISCUSSION

Species composition and characteristics

Streblus macrophyllus exhibited size structure with reverse J-shaped distribution suggesting continuous population regeneration and environmental suitability for establishment and survival of seedlings (Takahashi et al. 2001, Bunyavejchewin et al. 2003). There were many small trees which could potentially replace ageing adult trees and maintain the abundance of the species. This distribution type of diameter structure is often found for shade-tolerant species because one death of an adult will be replaced by individuals growing up from the smaller size classes (Richards 1996). Among the 40 remaining species, *C. tonkinensis* was a shade-intolerant species (FIPI 1996) with only 29 individuals but it contributed 41.79% of the total basal area.

Environmental homogeneity

Aggregation of individuals in plant communities at scales smaller than 20 m is often interpreted as seed dispersal limitation or facilitation (Hubbell et al. 2001, Uriarte et al. 2004) while at larger scales, it may be obscured by environmental heterogeneity, e.g. rock outcrops, slope or soil nutrients (Harms et al. 2001, John et al. 2007, Wiegand et al. 2007). Here, our hypothesis of environmental heterogeneity was tested based on the density variation of all adult trees at large scales by contrasting results from L-function and g-function. By means of Figure 4, the inference was confirmed that no large-scale aggregation was captured in tree density and, therefore, the hypothesis of a homogenous environment at our plot scale was accepted with approximate 99% confidence intervals. Moreover, this initial inference allowed us to select appropriate null models for further tests on ecological hypotheses of tree interaction or association in the study plot.

Overall univariate patterns

Aggregated distributions were found in both of the early life-history stages of *S. macrophyllus*, but the intensity and range of spatial scales notably decreased from juvenile to subadult stages. The aggregated patterns of juvenile and, to a smaller extent, subadult trees combined with the regular pattern of adult trees at small scales indicated that the spatial patterns of *S. macrophyllus* shifted with increasing tree sizes towards more regular patterns. This is evidence of density-dependent thinning that occurred within tree species, also known as self-thinning, which resulted from competition for natural resources including light, space, moisture, and nutrients (Kenkel 1988, Okuda et al. 1997). Similar patterns were found in other tropical forest trees (Sterner et al. 1986, Lan et al. 2009).

Contrasting spatial patterns of all species, *S. macrophyllus* and the remaining species together clearly showed the predominant effect of this species on patterning of the entire community. For the remaining species, juveniles and subadults showed marginally aggregated patterns while adult trees were randomly distributed. Significant patterns of all trees were presented again in *S. macrophyllus*. These findings confirmed the predominance and strong competitive interaction of *S. macrophyllus* beyond the dominance in abundance.

Dispersal of *S. macrophyllus*

We found lack of young trees in proximity to conspecific adults. This suggested density-dependent mortality because, due to seed rain, the density of seedlings was expected to be high in the vicinity of parent trees. Thereafter, it could be decreased by self-thinning. In the vicinity of *S. macrophyllus* adults, we found that juvenile and subadult trees occurred in a similar distribution as themselves. This is likely an outcome of the self-thinning process within *S. macrophyllus*. In addition, juveniles and subadults were clustered independently from adult trees. A potential mechanism that can produce the observed patterns of *S. macrophyllus* is a gap-phase regeneration mechanism (Hamill & Wright 1986, Condit et al. 1992). For example, if seedlings required more light to survive and recruit into juvenile and subadult stages, then the distributions of juveniles and subadult trees could be aggregated in less shaded sites or canopy gaps (Itoh et al. 1997). Figures 5 and 6 also suggested that density-dependent mortality (Kenkel 1988, Itoh et al. 1997, Condit et al. 2000) was an important process within *S. macrophyllus* leading to decreased numbers of young trees near potential parent trees as well as more regular distances between survivors. In addition, *S. macrophyllus* has fleshy fruits that may be dispersed away from mother trees by birds or mammals. Consequently, the aggregation of juveniles and subadults at scales up to 15 m suggested a patchy distribution at local scales or gap-phase regeneration mode (Grau 2000, Franklin & Santos 2011).

Heterospecific association of *S. macrophyllus*

The interspecific associations between adult trees of the remaining species (pattern 1) versus different life-history stages of *S. macrophyllus* (pattern 2) were expected to identify the local neighbourhood relationships of *S. macrophyllus* during its life-history. Interestingly, *S. macrophyllus* juveniles were weakly repulsed by other adults. Together with results from Figure 6, *S. macrophyllus* could be considered a light-demanding species at the juvenile stage. In the later life-history stages of *S. macrophyllus*, no significant association was found with the adults of other species (Figures 7a–b). This meant that *S. macrophyllus* adult trees competed strongly only with conspecific neighbours as shown in Figure 5 and were

independent of heterospecific adult neighbours. Therefore, light seemed to be a limiting factor for growth of *S. macrophyllus* in early life stages. From Figures 6 and 7, one potential explanation for the observed patterns was that *S. macrophyllus* might be an intermediate shade-tolerant species that needed sufficient sunlight to survive and grow in the early stages of life. Its patchy distribution seemed to be regulated by density-dependent mortality process.

CONCLUSIONS

Analyses of size structure of *S. macrophyllus* suggested that this species could continuously recruit and probably persisted as a population. Spatial pattern analyses via life-history stages showed evidence of self-thinning processes whereby recruitment into larger tree size seemed to depend on open canopy. Therefore, *S. macrophyllus* was characterised by a gap-phase regeneration mode. However, this study was based on a snap-shot inventory and limited by the plot size. Thus, some ecological dynamic processes such as recruitment, dispersal and mortality could not be captured in detail. Therefore, long-term measurements are required to deeper elucidate the ecological processes in spatio-temporal plant dynamics of this tree species.

ACKNOWLEDGEMENTS

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