Title: Large frugivores matter more on an island: insights from island-mainland comparison of

plant-frugivore communities

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ABSTRACT

Endozoochory, a mutualistic interaction between plants and frugivores, is one of the key

processes responsible for maintenance of tropical biodiversity. Islands, which have a smaller

subset of plants and frugivores when compared with mainland communities, offer an interesting

setting to understand the organization of plant-frugivore communities vis-a-vis the mainland

sites. We examined the relative influence of functional traits and phylogenetic relationships on

the plant-seed disperser interactions on an island and a mainland site. The island site allowed us

to investigate the organization of the plant-seed disperser community in the natural absence of

key frugivore groups (bulbuls and barbets) of Asian tropics. The endemic Narcondam Hornbill,

was the most abundant frugivore on the island and played a central role in the community.

Species strength, a measure of relevance of frugivores for plants, of frugivores was positively

associated with their abundance. Among plants, figs had the highest species strength and played

a central role in the community. Island-mainland comparison revealed that the island plant-seed

disperser community was more asymmetric, connected and nested as compared to the mainland

community. Neither phylogenetic relationships or functional traits (after controlling for

phylogenetic relationships) were able to explain the patterns of interactions between plants and

frugivores on the island or the mainland pointing towards the diffused nature of plant-frugivore

interactions. The diffused nature is a likely consequence of plasticity in foraging behavior and

trait convergence that contribute to governing the interactions between plants and frugivores.

Key words: community phylogenetics, insular communities, Narcondam Hornbill, plant-

frugivore interactions, seed dispersal

INTRODUCTION

Endozoochory, a mutualistic interaction between plants and frugivorous animals, represents a critical stage in plant regeneration and contributes to the maintenance of tree diversity in tropics (Terborgh et al. 2002). The community of fleshy-fruited plants and seed dispersers form a network of bipartite interactions. These networks could be influenced by different ecological (e.g. functional traits, phenological overlap) and evolutionary (e.g. co-evolution) processes (Vázquez et al. 2009; Guimarães Jr et al. 2011) whose relative roles are yet to be sufficiently investigated.

Shared evolutionary history can play a role in governing the interactions between plants and frugivores. Traits relevant to the seed dispersal such as beak width have been demonstrated to exhibit a phylogenetic signal (Rezende et al. 2007). On the other hand, trait convergence could result in unrelated species interacting with similar partners (Peralta 2016). Thus, interactions could be influenced by trait complementarity – which itself could be modulated by trait conservatism – or by trait convergence (Rezende et al. 2007; Bastazini et al. 2017).

To assess the relative roles of functional traits and their phylogenetic history in determining patterns of ecological interactions, Bazatazini et al. (2017) present an ecophylogenetic approach which outlines four scenarios based on the relationship between the degree of phylogenetic conservatism in species traits and their phylogenetic history. The four scenarios enable determining whether the interactions between plants and frugivores are 1) mediated by traits (with no trait conservatism), 2) mediated by phylogeny (with trait conservatism), 3) mediated by phylogeny (with no trait conservatism), or 4) mediated by both traits and phylogeny (with some trait conservatism) (see Fig. 1 in Bastazini et al. 2017 for additional details).

The influence of phylogenetic component would indicate the role of evolutionary processes like

co-evolution in the organization of the community, while the influence of functional component

controlling for phylogenetic history would imply trait complementarity. An integrated approach

that examines the role of evolutionary processes in tandem with ecological processes thus

enables an understanding of the relative roles of these factors in patterns of community

assembly. However, while the substantial phylogenetic data now available has improved our

understanding of community assemblages, the complementary empirical data that would allow

for an integrated approach has been lacking.

Despite zoochory being a pervasive interaction in the wet tropics of Asia, there is limited

information on plant-disperser communities from the mainland sites in the region (Corlett 2017)

and very poor information from oceanic islands in the Indian Ocean (Escribano-Avila et al.

2018). Most of the information on plant-disperser communities worldwide comes from mainland

sites with few studies from oceanic islands and even few studies set in an island-mainland

comparative framework (Schleuning et al. 2014).

Given this background, we examined the organization of the plant-disperser community on a

remote and small tropical island (Narcondam Island) in South Asia using the network and

ecophylogenetic approaches (Fig. 1A). The Narcondam island, which is the only place where the

Narcondam Hornbill *Rhyticeros narcondami* is found (Fig. 1B), is characterized by a species-

poor frugivore assemblage, which also enabled us to examine the relative quantitative role of

different species in seed dispersal.

In the Asian tropics, small-bodied frugivores like bulbuls and barbets are known to play key

roles in seed dispersal (Naniwadekar et al. 2019). However, the Narcondam Island does not have

any bulbuls and barbets and most of the dominant frugivores on the island are medium- and

large-bodied avian frugivores (> 100 g) (Fig. 1C). This offers an interesting system to understand

the pattern of interactions in absence of key frugivore group. We compared this island

community with a mainland community from north-east India with which it shares several

frugivore and plant species.

The specific aims of the study were to 1) identify key seed dispersers and plant groups on an

oceanic island community by comparing species-level properties (degree, species strength and

weighted betweenness) across different species, 2) examine the influence of abundance and

morphological traits on species strength of plants and frugivore species, 3) compare the network-

level properties (web asymmetry, connectance, specialization and weighted nestedness) of the

island community with a mainland community, and 4) examine the influence of phylogenetic and

functional component (fruit and seed size for plants and body size and degree of frugivory for

frugivores) on the distribution of interactions between plants and frugivores on the island and

mainland communities. This system allowed us to explicitly examine the change in the role of

large-bodied frugivores in plant-seed disperser communities in the absence of the important

small-bodied frugivores in Asian tropics (Fig. 1C). We hypothesized that large-bodied

frugivores, unhindered by morphological constraints, will demonstrate dietary shifts on islands in

the absence of small-bodied frugivores and will feed on fruits which they typically do not feed

on in the mainland site. This will alter the organization of the plant-seed disperser community on

the island as compared to mainland.

104　MATERIALS AND METHODS

Study area: The study was carried out on a small (area: 6.8 km^2), volcanic island, the

Narcondam Island (13º30'N, 94º38'E), in the Andaman Sea in India. The island is part of the

Indo-Myanmar Biodiversity Hotspot (Myers et al. 2000). Three distinct vegetation types on the

island include the narrow strip of littoral forests, forests dominated by deciduous species in the

north-eastern part of the island and evergreen forests in most of the island (Page et al. in prep.).

The Endangered Narcondam Hornbill *Rhyticeros narcondami* is found only on the Narcondam

Island (BirdLife International 2017). Around 50 species of birds have been reported from the

island (Raman et al. 2013). There are two terrestrial mammals on the island, including a shrew

(*Corcidura* sp.) and an invasive rat (*Rattus* sp.). There are at least four species of bats on the

island including a fruit bat species (*Pteropus hypomelanus*). Narcondam Hornbill (body mass:

600-750 g) is the largest frugivore on the island. Additional details of study area are in ESM

Online Resource 1.

Frugivore and plant abundance: We conducted the study between December 2019 – February

2020. We used variable-width line-transect surveys to estimate the density and abundance of

birds on Narcondam Island (Thomas et al. 2010). We walked 20 unique trails on the island (ESM

Online Resource 2: Fig. S1). We divided the entire elevation gradient into three elevation zones:

low (0-200 m), mid (200-400 m) and high (400-700 m) based on topography, vegetation

structure and composition. One to three observers walked trails in the mornings and afternoons.

During the trail walks, we recorded species identity, the number of individuals and the

perpendicular distance of the sighting to the trail. The trail lengths varied between 0.22–1.26 km

and the total sampling effort was 51.4 km (ESM Online Resource 2: Table S1).

We laid 49 belt transects (50 m × 10 m) across the three different elevation zones (low: 18, mid:

14, high: 17) to estimate the richness and abundance of woody plants on the Narcondam Island.

Within the belt, we enumerated all wood plants that were ≥ 10 cm GBH (girth at breast height)

along with their girth and taxonomic identity.

Plant-frugivore interactions: To systematically document plant-frugivore interactions, we

performed spot-sampling during our trail walks, and opportunistically outside our trail walks

following Palacio *et al*. (2016). Whenever we encountered a frugivore on a fruiting tree, we

recorded the species identity, number of frugivores, and frugivore foraging behaviour (fruits

swallowed, pecked or dropped) to ascertain whether the frugivore was a seed disperser or not.

Plant-frugivore interaction information was systematically collected outside trail sampling

whenever frugivore was detected on fruiting trees. To determine if we were missing

documenting interactions of frugivore species with low detection probability through spot

sampling, we systematically observed fruiting trees. We observed 40 fruiting individuals of 12

species following Naniwadekar et al. (2019a) (ESM Online Resource 2: Table S2). Additional

details of the tree watches are in ESM Online Resource 1. We measured widths of at least five

fruits each of 23 species of fleshy-fruited plants following Naniwadekar et al. (2019*a*) (ESM

Online Resource 2: Table S3). Plants were classified as small-seeded (seed width: < 5 mm),

medium-seeded (seed width: 5-15 mm) and large-seeded (> 15 mm) following Naniwadekar et

al. (2019a) (ESM Online Resource 2: Table S3).

Analysis

Frugivore and plant abundance: We estimated densities of perched birds with more than 40

visual detections using the R package 'Distance' (Thomas et al. 2010; Miller et al. 2019). We

estimated combined density of two Imperial-pigeon species (Pied Imperial-Pigeon *Ducula*

bicolor and Green Imperial-Pigeon *Ducula aenea*), since we had 45 detections overall.

Additional details of the analysis are provided in the ESM Online Resource 1. We estimated the

overall tree density and basal area ($m^2 ha^{-1}$), and density of 27 species of fleshy-fruited plants

that were fruiting during the study period.

Island plant-seed disperser community: We created a matrix of plants and frugivores with the

frequency of interactions documented for each plant-frugivore combination through spot

sampling. We only considered those animals as seed dispersers which we had seen swallowing

the seeds. We didn't consider Alexandrine Parakeet *Psittacula eupatria* in the analysis as we

documented the parakeets predating on seeds or feeding on unripe fruits as has also been

documented elsewhere (Shanahan et al. 2001). We plotted the interaction accumulation curve

using the 'random' method and 1000 permutations as implemented in the R package 'vegan' to

determine if we had adequately documented the diversity of unique plant-seed disperser

interactions (Oksanen et al. 2019).

To determine the relative roles of species in the organization of plant-seed disperser

communities, we examined species-level properties like the degree, species strength and

weighted betweenness centrality. Degree is defined as number of mutualistic partners of the focal

species, species strength is the sum of level of dependencies of all the partner species on the

focal species and betweenness centrality is a useful measure to identify connector species which

connects multiple guilds in a community (Dormann et al. 2008; Martín González et al. 2010).

We used the R package 'bipartite' for this analysis (Dormann et al. 2008).

We used the general linear model to examine the influence of plant abundance (estimated using

the vegetation plots) and fruit type (small-seeded, figs, medium-seeded and large-seeded) on

species strength of the 27 plant species that were fruiting during the study period. During

exploratory analysis, we noticed that figs consistently had higher species strength values. Since

certain fig groups are known to be keystone species in tropical forests and provide key nutrients

(Shanahan et al. 2001), we performed the analysis with figs as a separate category. Species

strength was $\log_{10}$ transformed to approximate normality. In the case of frugivores, since there

were only seven species and species strength values were not normally distributed, we used

Spearman's ρ to examine the correlation between abundance (encounter rate km^{-1}) of frugivores

and body mass with species strength. We obtained body mass information from Wilman et al.

(2014). To determine if the island plant-seed disperser community was organized into distinct

sub-communities, we used the program MODULAR (Marquitti et al. 2014) following

Naniwadekar *et al.* (2019). Additional details in ESM Online Resource 1.

Comparing the island and the mainland plant-seed disperser communities: To determine the

differences in the organization of plant-seed disperser communities between island and

mainland, we compared plant-disperser communities between the Narcondam Island with that of

Pakke Tiger Reserve in north-east India. Information on plant-seed disperser communities is

limited from the Asian tropics. However, there is detailed information on interactions between

47 avian frugivores and 43 plant species is available from Pakke Tiger Reserve in north-east

India (Naniwadekar et al. 2019). Pakke is part of the Eastern Himalaya Biodiversity Hotspot. It is

about 1500 km straight line distance north from the Narcondam Island. Unlike the island

community which is relatively young, the mainland site is several million years old.

Representative species of all the frugivore genera found on the Narcondam Island are also found

in Pakke with species like the Green Imperial-Pigeon, Asian Koel *Eudynamys scolopaceus* and

Common Hill Myna *Gracula religiosa* being reported from both sites. Several genera of fleshy-

fruited plants found on Narcondam Island are also found in Pakke. We examined network-level

properties like web asymmetry, connectance (proportion of realized interactions), specialization

(H_2') and nestedness (weighted NODF) on Narcondam Island and compared it with the mainland

site (Pakke). Values of web asymmetry range between -1 to 1 with negative values indicating

higher diversity of plant species as compared to animal species and vice versa (Blüthgen et al.

2007). Specialization is a measure of complementary specialization of the entire network with

high values indicating that the species are more selective. Weighted NODF is a quantitative

measure of nestedness with high values indicating higher nestedness.

Community assembly on island and mainland: To determine the influence of phylogeny and

functional traits on plant-frugivore interactions on the mainland and island site, we used the

following approach. The plant phylogeny was pruned from the megaphylogeny (Qian and Jin

2016) for both the island and mainland communities individually. This is a time-calibrated

phylogeny with largest representation from 98.6% of families and 51.6% of all genera of seed

plants. Scientific names of plants were standardized by referring to 'The Plant List'

(www.theplantlist.org). However, some species from Pakke and Narcondam were missing from

the megaphylogeny. In those cases, we chose the closest species from the Oriental region. When

we failed to find a congener, we chose a genus close to the focal genus (ESM Online Resource 2:

Table S3). We used fruit width and seed size class information for the analysis (ESM Online

Resource 2: Table S3). In the case of the birds, we referred to the comprehensive and time-

calibrated bird phylogeny available at www.birdtree.org (Jetz et al. 2012). We found all the focal

species for the two sites in the database except *Pycnonotus flaviventris*, *Chrysocolaptes*

guttacristatus and *Sitta cinnamoventris*. We used congeners from the Oriental region for these

species for the analysis (ESM Online Resource 2: Table S4). For the frugivores, we used body

mass and percentage fruits in the diet of frugivores (degree of frugivory) obtained from Wilman

et al. (2014) (ESM Online Resource 2: Table S4). We obtained 100 trees from posterior

distribution of 10000 trees based on Ericson Sequenced Species backbone phylogeny with 6670

OTUs for seven bird species for Narcondam and 47 species for Pakke found in the network (Jetz

et al. 2012). The maximum clade credibility tree was used for further analyses as these trees did

not vary substantially in topology and branch lengths. We calculated the phylogenetic distances

(pair-wise distances between species using their branch lengths) using 'cophenetic.phylo' in the

R package 'ape' (Paradis and Schliep 2018). We used Gower method to estimate pair-wise

distances among traits of both plants and frugivores using "vegdist" in the R package 'vegan'.

We then followed the novel approach outlined by Bastazini et al. (2017) to determine whether

phylogenetic proximity influenced the evolution of species trait and their interactions in the

network or whether species traits influence the observed pattern of interactions after removing

the potential confounding effect of the phylogeny. We used a weighted matrix of interactions

between plants and frugivores. It was frequency of interactions for the island site and visitation

rates for the mainland site. We applied this method for Narcondam and Pakke separately. We

used the R packages 'SYNCSA' for this analysis (Debastiani and Pillar 2012; Oksanen et al.

2019). We used R (ver. 3.5.3) for all the analysis (R Core Team 2019).

In addition, we assessed the phylogenetic signal separately in trait data for birds and plants in

both sites, Narcondam and Pakke. We calculated the Blomberg's *K* for each trait using the R

package "phytools" (Blomberg et al. 2003; Revell 2012).

RESULTS

Frugivore and plant abundance on island: We detected six avian frugivores during the trail

walks. We had 207 visual detections of Narcondam Hornbill, 71 detections of Asian Koel, and

45 detections of the two Imperial-Pigeons. We had only two detections each of the Common Hill

Myna and Eye-browed Thrush *Turdus obscurus*. Information on detection probability and mean

flock size is summarized in ESM Online Resource 2: Table S5.

Narcondam Hornbill was the most common frugivore on the island and its abundance (mean

(95% CI): 1026 birds (751-1402)) was almost four times that of the Asian Koel (Fig. 2A; ESM

Online Resource 1: Table S5). The estimated overall mean (95% CI) density of the Narcondam

Hornbill was 151 (110-206) birds km^{-2}. The mean densities of the Narcondam Hornbill varied

between 101-168 birds km^{-2} across the three elevation zones with the lowest mean densities in

the highest elevation band (Fig. 2B) (ESM Online Resource 2: Table S5).

The mean ($\pm$ SE) density and basal area of all the woody plants ($\geq$ 10 cm GBH) was highest in

the 0-200 m elevation band (density: 2091.1 $\pm$ 124.4 plants ha^{-1}; basal area (m^2 ha^{-1}): 58.3 $\pm$ 5.3),

followed by 200-400 m elevation band (density: 1505.7 $\pm$ 101.1 plants ha^{-1}; basal area (m^2 ha^{-1}):

49.1 $\pm$ 3.0) and least in the 400-700 m elevation band (density: 1311.8 $\pm$ 82.7 plants ha^{-1}; basal

area (m^2 ha^{-1}): 39.2 $\pm$ 4.2). However, the mean abundance of the different categories of the

frugivore food plant species (figs, small-seeded non-fig plants, medium-seeded plants and large-

seeded plants), which were fruiting during the study period, was highest in the high elevation

band (400-700 m), followed by middle elevation band and lowest in the low elevation band (0-

200 m) (Fig. 2C).

Island plant-seed disperser community: We documented 752 interactions between seven

frugivore species and 27 plant species during the spot scans performed on trail walks and

opportunistically. We documented all the frugivore species (except Andaman Green Pigeon

Treron chloropterus) previously reported from the island. We documented the migrant Daurian

Starling *Agropsar sturninus*, for the first time. During spot scans on trail walks, we documented

262 interactions between six frugivore species and 19 plant species. During opportunistic spot

scans, we documented 490 interactions between seven frugivore species and 26 plant species.

Narcondam Hornbill was documented feeding on 22 plant species, followed by Asian Koel,

which was documented feeding on 16 plant species (ESM Online Resource 2: Table S6). Of the

752 interactions, 533 were of the Narcondam Hornbill, 111 were of the Asian Koel, 36 of the

Green Imperial-Pigeon, 34 of the Pied Imperial Pigeon, 25 of the Common Hill Myna, 12 of the

Eye-browed Thrush and one of the Daurian Starling. Among the 27 plant species, eight species

were represented by *Ficus*. We observed 282 interactions on *Ficus rumphii*, 80 on *Ficus*

glaberrima, 54 on the small-seeded *Aidia densiflora* and 50 on the large-seeded *Caryota mitis*.

We documented 64 unique interactions between plants and frugivores on the island during the

study period. The interaction accumulation curve indicated that we had adequately documented

interaction diversity (ESM Online Resource 2: Fig. S2).

During tree watches, we documented 684 interactions (Narcondam Hornbill: 481, Asian Koel:

167, Pied Imperial-Pigeon: 23, Green Imperial-Pigeon: 7, Eye-browed Thrush: 5, Common Hill

Myna: 1), between 11 plant species and six frugivore species. We did not document any

interaction during tree watches which we had not documented during the spot scans. For eight of

the 11 tree species, we observed greater diversity of frugivores foraging on the particular plant

species during spot sampling as compared to tree watches.

Ficus rumphii (among plants) and Narcondam Hornbill (among frugivores) had the highest

degree, species strength and weighted betweenness centrality highlighting the central role they

played in the community during the study period (ESM Online Resource 2: Table S6). General

linear model indicated that the species strength in plants was influenced by plant type (figs and

non-fig small-, medium- and large-seeded plants) and not by plant abundance (Adj. $R^2 = 0.40$,

$F_{4,22} = 5.322$, $p = 0.004$) (Fig. 3A; ESM Online Resource 2: Table S7). *Ficus* were different from

other small-seeded plants and had higher species strength values (mean ($\pm$ SE) = 0.739 ($\pm$ 0.101),

n = 8 species) as compared to small-seeded (mean ($\pm$ SE) = 0.0816 ($\pm$ 0.01), n = 9 species),

medium-seeded (mean (± SE) = 0.0376 (± 0.01), n = 6 species) and large-seeded plant species

(mean (± SE) = 0.0328 (± 0.01); n = 4 species) (Fig. 3A). Among frugivores, species strength

was significantly and positively correlated to their abundance (Spearman's $\rho = 0.89$, $p = 0.0123$,

n = 7 species; Fig. 3B) and not significantly correlated to their body mass (Spearman's $\rho = 0.71$,

$p = 0.088$, n= 7).

The observed modularity of the island plant-disperser community was 0.33, and it was not

significantly different from that obtained through the null models ($p = 0.45$). The observed

community was subdivided into four modules and all the species were consistently allocated to

the same modules across the 25 runs (Table S8). One of the four modules comprised of the

Narcondam Hornbill and several medium and all the four large-seeded plants including

Endocomia macrocomia, *Caryota mitis*, *Pycnarrhaena lucida*, and *Planchonella longipetiolata*

highlighting hornbill's importance for large-seeded plants (Table S8).

Comparing the island and the mainland plant-seed disperser communities: The island

network was more asymmetric (-0.588) as compared to the mainland site (0.044). The observed

values of connectance and nestedness on Narcondam were almost twice that of the mainland site

and specialization (H_2') was almost half that of the mainland site (Table 1). The observed

network on Narcondam was less connected and less weighted than the null model predictions but

it was more specialized than expected by random (Table 1).

Community assembly on island and mainland: Among woody plants, we detected a strong

phylogenetic signal for seed size and fruit width in Pakke Tiger Reserve (seed size: Blomberg's

$K = 0.73$, $p = 0.001$; fruit width: Blomberg's $K = 0.43$, $p = 0.005$) and Narcondam (seed size:

Blomberg's $K = 1.48$, $p = 0.001$; fruit width: Blomberg's $K = 0.45$, $p = 0.027$). Among

frugivores, we detected a phylogenetic signal for the body mass of frugivores in Pakke

(Blomberg's K = 1.22, p = 0.012) and Narcondam Island (Blomberg's K = 1.28, p = 0.043). We

did not detect phylogenetic signal in the percentage of fruits in the diet of the frugivores of Pakke

(Blomberg's K = 0.23, p = 0.38) and Narcondam (Blomberg's K = 0.90, p = 0.08). We did not

find any significant correlation for the four models indicating that neither the functional nor the

phylogenetic component was able to explain the observed interaction patterns between plants and

frugivores in both island and mainland sites (Table 2; Fig. 4).

DISCUSSION

In this study, we demonstrate the role of large-bodied frugivores, especially the endemic

Narcondam Hornbill, in contributing disproportionately to seed dispersal in the absence of key

small-bodied frugivore groups on the island. Narcondam Hornbill plays a central role in the

plant-dispersal community as it is super-abundant, has very high species strength and

betweenness centrality values and is critical for the select large-seeded plants fruiting. This study

also demonstrates the key role played by figs in the plant-seed disperser community as

demonstrated by its very high species strength values as compared to other plant types. In

comparison to the mainland community, the island community was more asymmetric, connected,

nested and less-specialized. We have documented trait convergence and dietary shifts in

frugivores which potentially contribute to the diffused nature of plant-seed disperser interactions.

This also likely contributes to the absence of the role of phylogenetic or functional component in

the assembly of plant-seed disperser communities on mainland or the island. This is one of the

few studies from oceanic islands of the Indian Ocean and demonstrates how the largest

frugivore, which is a point-endemic, plays a critical quantitative role in seed dispersal. This is

also one of the few studies that performed a paired comparison between an island and mainland

community.

Narcondam for the hornbill and by the hornbill

There is no other place in the world where hornbills are reported to occur in such high densities

as on Narcondam Island (151 birds km^{-2}). On Sulawesi, the Knobbed Hornbills *Rhyticeros*

cassidix occur in very high densities of up to 84 birds km^{-2} (Kinnaird et al. 1996). In north-east

India, Wreathed Hornbill *Rhyticeros undulatus*, a close relative of the Narcondam Hornbill,

seasonally occur in densities of up to 68 birds km^{-2} (Naniwadekar and Datta 2013). Our density

estimates are similar to densities estimated by previous studies (Raman et al. 2013; Manchi

2017). Reduction in hunting pressures may have resulted in the recovery of Narcondam Hornbill

population from around 300 – 400 birds in early 2000s following the suggestions by Sankaran

(2000). Among other frugivores, there is limited information on the density of the Asian Koel

from other sites, but comparison with existing studies show that Narcondam has very high

densities of as compared to other sites (Riley 2001). The mean Imperial-Pigeon densities are

slightly lower than those reported from a mainland site in north-east India (Dasgupta and

Hilaluddin 2012).

In this study, multiple measures (abundance and different network metrics – species strength,

degree, betweenness and modularity) indicate that Narcondam Hornbills are keystone seed

dispersers on the island. Generally, as the body mass increases, the abundance of the frugivore

and the frequency of visitation is found to decline (Godínez-Alvarez et al. 2020). Interestingly,

the Narcondam Island is unique where the largest frugivore, the Narcondam Hornbill, is also the

most abundant frugivore on the island. Abundant species have been reported to influence their

partners strongly and Narcondam Hornbill had the highest species strength values highlighting

its importance for plants (Vázquez et al. 2007). Larger frugivores, like hornbills, are known to

remove more fruits per visit as compared to other smaller frugivores (Naniwadekar et al. 2019).

Highest number of interactions of the Narcondam Hornbill coupled with potentially higher fruit

removal rates point towards the key quantitative role played by the hornbill on the island.

Modularity analysis revealed that the hornbill was the key seed disperser for the large-seeded

plants as has been demonstrated in previous studies (Naniwadekar et al. 2019; Gopal et al. 2020).

The hornbill was the only frugivore that has high betweenness centrality value pointing towards

its vital role as a connector between different guilds within the community. This high reliance on

a frugivore and the potentially low redundancy highlights the fragility of the island ecosystem

because of strong reliance on a single frugivore.

Narcondam has very high densities of hornbill food plants, which likely explains the high

hornbill densities. The overall fig density on Narcondam Island was 27 trees ha^{-1}. This is very

high as compared to other sites in the world where the densities of hemi-epiphytic figs are

typically less than one tree ha^{-1} (Harrison 2005). *Ficus* (overall) densities were 1.4 trees ha^{-1} at a

site in the southern Western Ghats (Page and Shanker 2020). *Ficus* densities of up to 10 trees ha^{-}

^{1} have been reported in Sulawesi (Kinnaird et al. 1996). Consistently across the mainland site

and the island site, certain species of *Ficus* attracted the highest diversity of frugivores. On the

island, *Ficus* represented the top six species in species strength, highlighting the key role they

potentially played in the plant-disperser community at least during the study period. Association

between canopy, hemi-epiphytic figs and hornbills has been documented elsewhere (Harrison et

al. 2003). Most dominant fig species during our study on Narcondam Island, which are important

hornbill food plants are mostly canopy, hemi-epiphytic figs. Being small-seeded, they often play

the pioneering role in attracting frugivores on oceanic islands and accelerating plant colonization

on the island (Harrison 2005). Figs and hornbills may have played a critical role in setting up the

plant community on the island which needs further investigation.

Based on published literature on hornbill food plants from north-east (Datta 2001; Naniwadekar

et al. 2015, 2019) and Thailand (Poonswad et al. 1998; Kanwatanakid-Savini et al. 2009;

Chaisuriyanun et al. 2011), we identified at least 35 species of Narcondam Hornbill food plants.

Overall density of the 35 hornbill food plant species (GBH $\geq$ 30 cm) on Narcondam Island is 327

trees ha^{-1}. In Pakke Tiger Reserve, the density of 45 hornbill food plant species was 163 trees ha^{-}

^{1} (Datta 2001), highlighting the hyper abundance of hornbill food plants on Narcondam Island.

Interestingly, the abundance of food plants is consistently distributed across the entire elevation

gradient. The hyper abundance of frugivores and their food plants makes this island a unique

ecosystem.

Plant-seed disperser communities on island and mainland

Island communities have been reported to be more asymmetric as compared to mainland systems

due to higher immigration rates of plants or human-caused frugivore extinctions on the island

(Schleuning et al. 2014; Nogales et al. 2016). There are no known human-driven extinctions on

the Narcondam island. We are unlikely to find additional species of frugivores that can play an

important functional role in seed dispersal on the island as our list of frugivores is comparable to

the frugivores reported by the previous studies (Raman et al. 2013). We feel that dispersal

limitation of frugivores could also be a key factor in the web asymmetry. Andaman and Nicobar

archipelago of which Narcondam Island is part of, is an interesting example of this. Bulbuls and

barbets, which are the key frugivores on the mainland (Naniwadekar et al. 2019), are absent on

the remote and tiny Narcondam Island. Endemic species of bulbuls are found on other islands in

the Andaman and Nicobar archipelago, but barbets are absent from the archipelago. Thus, the

variation in the dispersal ability of different frugivore groups might also play an important role in

contributing towards the web asymmetry on oceanic islands. Among other network properties,

higher connectance on islands has been reported elsewhere (González-Castro et al. 2012).

Forbidden links are thought to contribute towards lower connectance (Olesen et al. 2011).

Additionally, connectance and nestedness are negatively correlated (Rezende et al. 2007). In our

case, higher connectance is associated with higher nestedness on the island site as compared to

the mainland site. Given that most of the important frugivores on the island are medium (100-

500 g) or large-bodied (> 500 g) with ability to handle small and large fruits (thereby less

likelihood of forbidden links), it has likely resulted in higher connectance and nestedness.

Organization of plant-seed disperser communities

We detected contrasting phylogenetic signals in bird body mass, and seed and fruit traits on both

the island and the mainland. While frugivore traits were more likely to be similar (than Brownian

motion model) for closely related species, plant traits were less likely to be similar for closely

related species. However, functional and phylogenetic component did not influence the network

interactions in the plant-seed disperser communities on the island and mainland. The role of

evolutionary history in shaping local networks of interacting species is thought to be limited

(Segar et al. 2020). Unlike the plant-pollinator networks, the seed dispersal networks can be

expected to be less specialized and diffused as unrelated species may exhibit trait convergence

and forage on similar plant species (Jordano 2000). We found that distantly related groups of

frugivores dispersed seeds of a common assemblage of plants, many of which are distantly

related to each other. A diverse array of frugivores disperse the super-generalist *Ficus*, and even

the large-seeded plants, which otherwise have a smaller disperser assemblage, were

predominantly dispersed by distant frugivore clades, like hornbills and *Ducula* pigeons, on the

mainland site. Unlike Bastazini et al. (2017), we did not detect signal in the functional

component of the plant-seed disperser communities. This is a likely consequence of the absence

of trait complementarity among interacting frugivores and plants. While the small-bodied

frugivores may not be able to disperse large seeds due to morphological constraints, large-bodied

frugivores like hornbills, fed on plants that were large- and small-seeded resulting in lack of trait

complementarity.

Another important reason could be the behavioral plasticity shown by frugivorous birds while

foraging. In the mainland site, where there are almost 50 frugivore species with a higher

preponderance of small-bodied frugivores, we rarely encountered *Ducula* pigeons on figs that

attracted more than half of the frugivore species found in the area. On the mainland site, *Ducula*

pigeons were mostly observed foraging on medium-seeded and large-seeded plants. On the

island, where there are hardly any small frugivores, we mostly saw the *Ducula* pigeons foraging

on *Ficus*. This behavior of *Ducula* pigeons feeding on figs on the island but less frequently on

the mainland demonstrates phenotypic plasticity in *Ducula* pigeons, likely as a response to lack

of competitive interactions, an aspect that has been seldom documented elsewhere. Additionally,

on Narcondam Island, we documented hornbills feeding on *Callicarpa arborea*, a small-seeded

species which is mostly dispersed by bulbuls and barbets on the mainland site (Naniwadekar et

al. 2019). Asian Koel, which is a rare frugivore in the mainland site, was seen only once feeding

on fruit there. However, on the island, it is the second-most important frugivore. It remains to be

determined, whether exploitative competition on the mainland site drives larger frugivores to

feed on different resources in the mainland forested system. While habitat fragmentation often

results in loss of large-bodied frugivores thereby affecting large-seeded plants, it will be

interesting to investigate the implications of the absence of small-bodied frugivore on the

structuring of plant communities.

Bats are known to play an important role in seed dispersal on oceanic islands (McConkey and

Drake 2015). We recorded spat out wads of pulp of three *Ficus* spp., *Balakata*, and

Planchonella. Direct detections of foraging bats on fruiting *Manilkara zapota* (planted),

Terminalia catappa, *Ficus rumphii* and *Ardisia*. We failed to detect foraging bats during the six

night-walks that lasted for at least 90 minutes each.

One may argue about the lack of phenological completeness of the study and its influence on the

inferences drawn. The abundances of the frugivores is unlikely to change since it is a small and

remote island in the middle of the Andaman Sea. Narcondam Hornbill is likely to remain the

most abundant frugivore on the island year-round and thereby play a key role in seed dispersal

throughout the year. Additionally, the frugivore species composition on the island is unlikely to

change as previous studies have documented similar assemblage of frugivores. We have a

diverse representation of plant families in our study with varying fruit (figs, drupes and arillate

dehiscent capsular fruits) and seed traits thereby capturing the functional diversity in traits of

fruits and seeds making the data amenable for ecophylogenetic analysis. Additionally, this is a

remote island making it extremely challenging to conduct long bouts of fieldwork, thereby

posing challenges to collect long-term data.

Only those plants and animals that are able to cross the oceanic barriers and establish contribute

to structuring of communities on islands. Degree of isolation, area and age of islands thereby

offers an interesting gradient to understand the relative importance of processes in structuring of

communities. While, we did not detect a phylogenetic or functional signal across mainland and

island communities, it will be interesting to investigate these patterns as the diversity of

frugivores increases further. Stronger signals can be expected in a niche packing scenario. As

demonstrated in this study, large-bodied frugivores, particularly the Narcondam Hornbill, play a

pivotal role on Narcondam Island in the absence of small frugivores on a relatively young and

moderately isolated island. Several threatened hornbill species are found on oceanic islands

(IUCN 2019). Narcondam Island and its endemic hornbill give insights into the potential role of

hornbills, which are among the largest frugivores on islands, in shaping and potentially

maintaining the island tree communities by playing a key role as a seed disperser. The ecological

role of hornbills and their contribution to mutualism and maintenance of plant species diversity is

poorly understood from other oceanic islands. Narcondam Island through the hyper abundance of

hornbills and its food plants and the asymmetric nature of the plant-disperser community with

heavy reliance on a single frugivore demonstrates the unique ecological and evolutionary setting

of oceanic islands. The disproportionate and singular role that species can play on oceanic

islands points towards the extreme vulnerability of oceanic islands to any external perturbations.

The perturbations can disrupt the delicate and intricate relationships between plants and seed

dispersers, with dramatic cascading impacts on the entire community. Unfortunately, studies like

this might not be possible in many other oceanic islands that harbor hornbills as they have been

already modified extensively due to anthropogenic activities. It is indeed fortunate that

Narcondam Island has been preserved as a Wildlife Sanctuary thereby safeguarding not only a

unique hornbill species but also a unique ecosystem.

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Table 1. Network-level properties (observed and null model values) for the Narcondam Island,

and observed values for the mainland site, Pakke Tiger Reserve. Null model values for the

mainland site can be found in Naniwadekar et al. (2019a).

	Observed value: Narcondam	Null mean: Narcondam	Null SD: Narcondam	Observed value: Mainland
Connectance	0.339	0.463	0.018	0.154
Specialization (H_2')	0.233	0.144	0.009	0.500
Weighted NODF	45.178	59.76	3.834	24.885

671   **Table 2.** Correlation coefficients for the two models that evaluate the influence of the functional

672   (after removing phylogenetic signal) and phylogenetic components on the observed patterns of

673   interactions between plants and frugivores for the mainland site (Pakke Tiger Reserve) and the

674   island site (Narcondam Wildlife Sanctuary). None of the values were significant at $p = 0.05$.

Parameter	ρ (Mainland)	ρ (Island)
Functional component (after removing phylogenetic signal)	0.04	0.03
Phylogenetic component	0.023	-0.17

FIGURES

Figure 1. Schematic illustrating mainland and island plant and seed-dispersal communities. a)

map of the South and Southeast Asia showing mainland and island site, b) the island endemic

large-bodied frugivore Narcondam Hornbill *Rhyticeros narcondami* on *Caryota mitis*, c)

Representative schematic showing organization of plant-seed disperser community on mainland

and island. Mainland community is more diverse and symmetric as compared to the island

community. On mainland, due to the presence of great diversity of small-bodied frugivores

which are more efficient in removal of small fruits, large bodied frugivores may predominantly

feed on large-seeded plants. On the island, in the absence of small-bodied frugivores, large-

bodied frugivores will start feeding on small fruits resulting in dietary shifts and more dominant

role in the community as indicated by the thickness of arrows. Dietary changes will in turn result

in changes in interaction frequencies with corresponding changes in network structure. Painting

and sketches by Sartaj Ghuman. Map source: Google Earth.

Figure 2. Abundance of Narcondam Hornbill *Rhyticeros narcondami*, Asian Koel *Eudynamys*

scolopaceus and Imperial-Pigeons (Pied Imperial-Pigeon *Ducula bicolor* and Green Imperial-

Pigeon *Ducula aenea*) on Narcondam Island (A). Density (km^{-1}) of Narcondam Hornbill on the

island and across the three elevation zones on the island (B). Density (ha^{-1}) of 27 species of

fleshy-fruited plants that were fruiting across the three elevation zones during the study period

(C).

Figure 3. Relationship between species strength (log_{10} transformed in the case of plants) and

plant type (A) and frugivore abundance (B). We separated figs and small-seeded plants for they

consistently had higher species strength values. Frugivore codes - ASKO: Asian Koel, CHMY:

Common Hill Myna, DAST: Daurian Starling, EBTH: Eye-browed Thrush, GIPI: Green

Imperial-Pigeon, NAHO: Narcondam Hornbill and PIPI: Pied Imperial-Pigeon.

Figure 4. Seed dispersal interactions in Narcondam Island (A) and the mainland site, Pakke

Tiger Reserve (B). Also shown is the pruned phylogeny for the plants and the frugivores and

respective traits for plants (seed size and fruit width) and frugivores (body mass and percentage

diet that is comprised of fruits). The matrix is weighted in both the island and mainland case and

it has frequency of interactions for the island site and visitation rate of frugivores (h^{-1}) for the

mainland site shown by the grey bar below the matrix. * indicates species for which a congener

or allied genus was used in the constructing the phylogenetic tree.

Figure 1.

709   **Figure 2.**

712   **Figure 3.**

715 **Figure 4.**

39