# **RESEARCH ARTICLE**

# Interspecific Interactions between Primates, Birds, Bats, and Squirrels May Affect Community Composition on Borneo

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For several decades, primatologists have been interested in understanding how sympatric primate species are able to coexist. Most of our understanding of primate community ecology derives from the assumption that these animals interact predominantly with other primates. In this study, we investigate to what extent multiple community assembly hypotheses consistent with this assumption are supported when tested with communities of primates in isolation versus with communities of primates, birds, bats, and squirrels together. We focus on vertebrate communities on the island of Borneo, where we examine the determinants of presence or absence of species, and how these communities are structured. We test for checkerboard distributions, guild proportionality, and Fox's assembly rule for favored states, and predict that statistical signals reflecting interactions between ecologically similar species will be stronger when nonprimate taxa are included in analyses. We found strong support for checkerboard distributions in several communities, particularly when taxonomic groups were combined, and after controlling for habitat effects. We found evidence of guild proportionality in some communities, but did not find significant support for Fox's assembly rule in any of the communities examined. These results demonstrate the presence of vertebrate community structure that is ecologically determined rather than randomly generated, which is a finding consistent with the interpretation that interactions within and between these taxonomic groups may have shaped species composition in these communities. This research highlights the importance of considering the broader vertebrate communities with which primates co-occur, and so we urge primatologists to explicitly consider nonprimate taxa in the study of primate ecology. Am. J. Primatol. 75:170-185, 2013. © 2012 Wiley Periodicals, Inc.

## Key words: niche differentiation; community assembly; Sciuridae; Chiroptera; southeast Asia

## **INTRODUCTION**

For several decades, primatologists have been interested in the study of primate communities [Bourliere, 1984; Fleagle & Reed, 1996; Reed & Bidner, 2004]. A major component of studying primate communities has been to understand how species within communities are able to coexist over time. According to the principle of competitive exclusion, no two species can survive on the same limiting resource [Gause, 1934] and as a consequence no two species can occupy the same niche [Hutchinson, 1957]. A great deal of effort has therefore gone into understanding how niche space is divided among primate species within communities. Niche differentiation among primates has arguably led to differences in diet, canopy use, habitat use, and body size among other dimensions [Bourliere, 1984; Fleagle et al., 1999]. In this research, primatologists have viewed primate communities as self-contained, and as a consequence, most of our understanding of primate

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community ecology derives from the assumption that primates interact predominantly with other primates. However, the full set of species that compete with primates likely includes various nonprimate taxa [Emmons et al., 1983; Estrada & Coates-Estrada, 1985; French & Smith, 2005; Ganzhorn, 1999; Gautier-Hion et al., 1980; Marshall et al., 2009b; Strum & Western, 1982].

The extent to which primates may compete with other taxonomic groups likely varies across sites and biogeographic regions. Indeed there is substantial variation across continents in the proportion of mammals that are primates [Thalmann, 2007]. Intriguingly, primates compose 44% of nonvolant terrestrial mammalian species in Madagascar, whereas in other regions primates constitute only 8-12% of nonvolant land mammals [Jernvall & Wright, 1998]. This difference is particularly interesting in light of recent studies that are consistent with the interpretation that competition between primates is stronger in Madagascar than other regions [Beaudrot & Marshall, 2011; Kamilar & Guidi, 2010]. Given that primates comprise a larger portion of the vertebrate community in Madagascar, these animals may be competing more with each other on this island than in other regions where primates may be competing more with other animals. If competition for food resources is more severe between primates and other animals than among primates, then the effects of competition on community structure may be masked if primates are studied in isolation. Biogeographic regions other than Madagascar are therefore strong candidates for investigating the role of competition with nonprimate taxa in shaping primate communities.

Community ecologists have developed a suite of methods to examine interspecific interactions at the community level. These include metrics for quantifying patterns of species distributions expected to occur if competition and other interactions strongly affect the presence and absence of species across sites, or in other words, if species interactions structure patterns of species co-occurrence. Over the past few decades, there has been considerable advancement in the statistical rigor with which co-occurrence patterns are analyzed [Gotelli & Entsminger, 2011; Stone & Roberts, 1990]. Moreover, much research has incorporated these metrics to evaluate the importance of interspecific interactions in structuring communities, including a meta-analysis of 96 communities in which the strongest results were found for mammals and birds [Gotelli & McCabe, 2002]. Similar patterns occur in microorganisms, which suggests that co-occurrence patterns shaped by interspecific interactions may be present across a wide variety of life forms [Horner-Devine et al., 2007].

In this study, we investigated community assembly hypotheses for primate species in isolation and in addition to a larger community of vertebrates whose arboreal foraging and diet suggest a competitive relationship with sympatric primates. We focused on vertebrate communities in forested areas on Borneo, which is highly species-rich and is the world's second largest island. Of the biogeographic regions in which living primates occur, Asian primate communities are characterized by low species richness relative to other taxonomic groups and also in comparison to primate communities of other regions [Reed & Bidner, 2004; Thalmann, 2007]. Moreover, Ganzhorn [1999] has argued that competition between primates and other taxa is stronger on Borneo than in other regions based on analyses of body mass ratios. We examined to what extent communities are shaped by interspecific interactions when we examined communities of primates alone and in combination with other taxonomic groups. Specifically, we investigated three tests for detecting the effects of species interactions on communities: checkerboard distributions, guild proportionality, and Fox's assembly rule for favored states.

## **Checkerboard Distributions**

One approach to evaluating species interactions is to look for evidence of significant patterning in the presence or absence of species across communities that has resulted from interactions between pairs of species. For example, a checkerboard distribution refers to the alternating presence of ecologically similar species on islands or other habitat patches, such as protected areas. In a checkerboard distribution, species A is present but species B is absent in some patches while species B is present but species A is absent in other patches (Fig. 1). Although there has been a long history in ecology of interpreting significantly checkerboard results as evidence of competition exclusion between pairs of species [Diamond, 1975; Gotelli & McCabe, 2002; Horner-Devine et al., 2007; Stone & Roberts, 1990], recent modeling efforts have suggested that interspecific interactions other than competition may produce significant checkerboard results [Ulrich & Gotelli, 2012]. Although interspecific competition is one

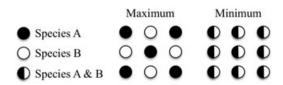


Fig. 1. Illustration of the maximum and minimum extent of a checkerboard distribution of two species. Each circle represents a site that contains either species A shown in black, species B shown in white, or both species. If species A and B are not found at the same site, then they exhibit a maximally checkerboarded distribution, whereas if species A and B consistently co-occur at the same site then they exhibit a minimally checkerboarded distribution.

potential causal factor driving checkerboard distributions, alternative explanations such as habitat checkerboard and historical effects are also possible and warrant additional consideration if analyses produce significant checkerboard results [Gotelli & McCabe, 2002]. We therefore investigated checkerboard distributions to evaluate the presence or absence of significant ecological structure in primate communities alone versus in the context of the broader vertebrate communities and suggest that results can direct future research towards understanding causal mechanisms.

# **Guild Proportionality**

A guild refers to a group of species consuming the same class of resources [Fauth et al., 1996]. If competition affects species within communities at the guild level rather than at the level of individual species (i.e. if species within a guild are equivalent and guild-level resources are critical to determining the number of species that can persist in a guild), then the proportion of species in a guild should be relatively constant across communities irrespective of the identity of the species [Wilson, 1989]. For example, if we consistently find that each community consists of 50% frugivores, 30% insectivores, 20% omnivores, then the percentage of species in each guild is proportional across communities. Guild proportionality derives from the assumption that resource distribution is proportional across communities and because it is based strictly on the proportion of species, it does not take into account differences in species' body sizes that may affect resource use. There is evidence of guild proportionality in avian, ant, grassland, and saltmarsh communities [Feeley, 2003; Holdaway & Sparrow, 2006; Silva & Brandao, 2010; Wilson & Whittaker, 1995], but to our knowledge previous tests have not investigated mammals.

## Fox's Assembly Rule for Favored States

According to Fox's assembly rule [Fox, 1987], interspecific competition should favor the entry and persistence of species that are most distinct from those already present in a community. The rule predicts that an incoming species will more likely persist if it belongs to a guild not represented in the community than if it attempts to divide niche space with species of the same guild that are already present. If interspecific competition favors the entry of species that are most distinct from members already present in the community, then communities will contain an equal number of species in each guild more often than is expected by chance. The rule defines a site as favored (i.e. in support of the assembly rule) if and only if species fill each guild evenly such that no guild has two or more representatives than another guild. For example, according to the rule a site with three frugivores, three omnivores, and four insectivores is favored whereas a site with three frugivores, three omnivores, and five insectivores is unfavored.

Fox's assembly rule is similar to guild proportionality in that it assumes that the distribution of resources is even across sites, however it also requires the more strict assumption that resource distribution is even within sites. Hence, Fox's assembly rule assumes that the distribution of all resources is homogeneous and therefore that the number of species at each site in each guild should be more homogeneous than expected by chance. For a group of sites, if more sites meet the test for "favored" than are expected based on a null model, then the sites support the assembly rule.

# **METHODS**

# **Data Collection**

We compiled species presence–absence data on bats, birds, primates, and squirrels (Table I) from a combination of published and unpublished sources for 21 forested sites located throughout Borneo [Struebig et al., 2010]. We included as many sites for which we could obtain reliable species lists of all four taxonomic groups. The sites include all major naturally occurring habitat types found in Borneo (i.e. mangrove, swamp, lowland, hill, montane, and karst forests). For a subset of the analyses, we restricted the number of sites to include only those with the same broad habitat types to control for variation in habitat. These sites included lowland, hill, and montane forest (Fig. 2).

We used published descriptions to classify each species based on arboreality (terrestrial or nonterrestrial), activity pattern (diurnal or nocturnal), resident status (resident or nonresident), and diet (folivore, frugivore, frugivore/folivore, insectivore, or omnivore;). We included only resident, arboreal species because these species are most likely to interact with primates. To assign species to guilds, we used broad dietary categories to code for the presence or absence of items in the diet of each species based on qualitative diet descriptions. We defined folivores as species whose diets contained consistently available items, such as vegetation, moss, lichen, fungus, bark, or roots. We defined frugivorous species as those species for which the diet contains fruit, nectar, seeds, buds, or flowers. We designated species with dietary items in both of these categories as frugivore/folivores. Insectivores included species that exclusively fed on insects or other invertebrates. Omnivores included species that fed on both plant and animal matter. We excluded any species with a diet containing vertebrates because the diet would unlikely overlap with primates unless the diet also contained plant

Class	Order	Family	Genus	Species	No. of Sites	Guild	Invert	Fruit	Act
Aves	Accipitriformes	Accipitridae	Aviceda	jerdoni	7	NA	0	0	D
Aves	Apodiformes	Apodidae	Aerodramus	fuciphagus	9	IN	1	0	D
Aves	Apodiformes	Apodidae	Aerodramus	maximus	10	IN	1	0	D
Aves	Apodiformes	Apodidae	Aerodramus	salangana	$\frac{2}{9}$	IN	1	0	D
Aves	Apodiformes Apodiformes	Apodidae Apodidae	Apus Collocalia	affinis esculenta	9 10	IN IN	1 1	0 0	D D
Aves Aves	Apodiformes	Apodidae	Conocana Cypsiurus	balasiensis	10	IN IN	1	0	D
Aves	Apodiformes	Apodidae	Hirundapus	gigateus	4 9	IN	1	0	D
Aves	Apodiformes	Apodidae	Hydrochorous	gigas	3	IN	1	0	D
Aves	Apodiformes	Apodidae	Rhaphidura	leucopygialis	11	IN	1	Ő	D
Aves	Apodiformes	Hemiprocnidae	Hemiprocne	comata	14	IN	1	0	D
Aves	Apodiformes	Hemiprocnidae	Hemiprocne	longipennis	14	IN	1	0	D
Aves	Bucerotiformes	Bucerotidae	Anthracoceros	albirostris	8	OM	1	1	D
Aves	Bucerotiformes	Bucerotidae	Anthracoceros	malayanus	14	OM	1	1	D
Aves	Bucerotiformes	Bucerotidae	Berenicornis	comatus	11	OM	1	1	D
Aves	Bucerotiformes	Bucerotidae	Buceros	rhinoceros	16	OM	1	1	D
Aves	Bucerotiformes	Bucerotidae	Rhinoplax	vigil	17	OM	0	1	D
Aves	Bucerotiformes	Bucerotidae	Rhyticeros	corrugatus	15	OM	1	1	D
Aves	Bucerotiformes	Bucerotidae	Rhyticeros	undulatus	11	OM	1	1	D
Aves	Caprimulgiformes	Caprimulgidae	Caprimulgus	affinis	1	IN	1	0	N
Aves	Caprimulgiformes Columbiformes	Caprimulgidae Columbidae	Caprimulgus	concretus	$\frac{3}{7}$	IN FR	1 0	$\begin{array}{c} 0 \\ 1 \end{array}$	N D
Aves Aves	Columbiformes	Columbidae	Ducula Ducula	aenea badia	6	FR	0	1	D
Aves	Columbiformes	Columbidae	Ducula	bicolor	5	FR	0	1	D
Aves	Columbiformes	Columbidae	Ducula	pickeringii	$\frac{5}{2}$	FR	0	1	D
Aves	Columbiformes	Columbidae	Macropygia	emiliana	$\frac{2}{5}$	FR	0	1	D
Aves	Columbiformes	Columbidae	Macropygia	ruficeps	$\tilde{5}$	FR	Ő	1	D
Aves	Columbiformes	Columbidae	Ptilinopus	jambu	8	FR	0	1	D
Aves	Columbiformes	Columbidae	Ptilinopus	melanospilus	1	$\mathbf{FR}$	0	1	D
Aves	Columbiformes	Columbidae	Treron	capellei	14	$\mathbf{FR}$	0	1	D
Aves	Columbiformes	Columbidae	Treron	curvirostra	12	$\mathbf{FR}$	0	1	D
Aves	Columbiformes	Columbidae	Treron	fulvicollis	4	$\mathbf{FR}$	0	1	D
Aves	Columbiformes	Columbidae	Treron	olax	10	$\mathbf{FR}$	0	1	D
Aves	Columbiformes	Columbidae	Treron	vernans	7	$\mathbf{FR}$	0	1	D
Aves	Coraciiormes	Meropidae	Nyctyornis	amictus	14	IN	1	0	D
Aves	Cuculiformes	Cuculidae	Cacomantis	merulinus	11	OM	1	1	D
Aves	Cuculiformes Cuculiformes	Cuculidae Cuculidae	Cacomantis Cacomantis	sonneratii variolosus	8 4	IN IN	1 1	0 0	D D
Aves Aves	Cuculiformes	Cuculidae	Chrysococcyx	minutillus	4	IN IN	1	0	D
Aves	Cuculiformes	Cuculidae	Chrysococcyx	russatus	1	IN	1	0	D
Aves	Cuculiformes	Cuculidae	Chrysococcyx	xanthorhynchus	8	OM	1	1	D
Aves	Cuculiformes	Cuculidae	Hierococcyx	fugax	12	OM	1	1	D
Aves	Cuculiformes	Cuculidae	Hierococcyx	vagans	6	IN	1	0	D
Aves	Cuculiformes	Cuculidae	Phaenicophaeus (Rhopodytes)	diardi	11	IN	1	0	D
Aves	Cuculiformes	Cuculidae	Rhinortha	chlorophaea	13	IN	1	0	D
Aves	Cuculiformes	Cuculidae	Surniculus	lugubris	15	OM	1	1	D
Aves	Cuculiformes	Cuculidae	Zanclostomus	javanicus	11	IN	1	0	D
Aves	Falconiformes	Falconidae	Microhierax	latifrons	3	NA	0	0	D
Aves	Passeriformes	Acanthizidae	Gerygone	sulphurea	11	IN	1	0	D
Aves	Passeriformes	Aegithinidae	Aegithina	tiphia	7	IN	1	0	D
Aves	Passeriformes	Aegithinidae	Aegithina	viridissima Gun hainta	13	IN	1	0	D
Aves	Passeriformes	Campephagidae	Coracina Coracina	fimbriata Iamata	13	OM OM	1	1 1	D D
Aves	Passeriformes Passeriformes	Campephagidae	Coracina Coracina	larvata striata	5 9	ОМ IN	1 1	1	D D
Aves	Passeriformes	Campephagidae Campephagidae	Coracina Hemipus	striata hirundinaceus	9 15	IN IN	1		D
Aves Aves	Passeriformes	Campephagidae	Hemipus Hemipus	nirunainaceus picatus	15 8	IN IN	1	0	D
Aves	Passeriformes	Campephagidae	Lalage	nigra	8 4	IN IN	1	0	D
Aves	Passeriformes	Campephagidae	Pericrocotus	flammeus (speciosus)	14	IN	1	0	D

# TABLE I. List of species included in the study

Class	Order	Family	Genus	Species	No. of Sites	Guild	Invert	Fruit	Act
Aves	Passeriformes	Campephagidae	Pericrocotus	igneus	10	IN	1	0	D
Aves	Passeriformes	Campephagidae	Pericrocotus	solaris	2	IN	1	0	D
Aves	Passeriformes	Chloropseidae	Chloropsis	cochinchinensis	13	OM	0	0	D
Aves	Passeriformes	Chloropseidae	Chloropsis	cyanopogon	16	OM	0	0	D
Aves	Passeriformes	Chloropseidae	Chloropsis	sonnerati	15	OM	1	1	D
Aves	Passeriformes	Corvidae	Cissa	thalassina	1	IN	1	0	D
Aves	Passeriformes	Corvidae	Dendrocitta	cinerascens	4	OM	1	1	D
Aves	Passeriformes	Corvidae	Platylophus	galericulatus	13	IN	1	0	D
Aves	Passeriformes	Corvidae	Platysmurus	leucopterus (aterrimus)	11	IN	1	0	D
Aves	Passeriformes	Dicaeidae	Dicaeum	agile	2	OM	1	1	D
Aves	Passeriformes	Dicaeidae	Dicaeum	chrysorrheum	12	OM	1	1	D
Aves	Passeriformes	Dicaeidae	Dicaeum	concolor (minullum)	10	OM	1	1	D
Aves	Passeriformes	Dicaeidae	Dicaeum	cruentatum	11	OM	1	1	D
Aves	Passeriformes	Dicaeidae	Dicaeum	everetti	4	OM	1	1	D
Aves	Passeriformes	Dicaeidae	Dicaeum	monticolum	6	OM	1	1	D
Aves	Passeriformes	Dicaeidae	Dicaeum	trigonostigma	13	OM	1	1	D
Aves	Passeriformes	Dicaeidae	Dicaeum	trochileum	3	OM	1	1	D
Aves	Passeriformes	Dicaeidae	Prionochilus	maculatus	15	$\mathbf{FR}$	1	0	D
Aves	Passeriformes	Dicaeidae	Prionochilus	percussus	7	$\mathbf{FR}$	1	0	D
Aves	Passeriformes	Dicaeidae	Prionochilus	thoracicus	11	OM	1	1	D
Aves	Passeriformes	Dicaeidae	Prionochilus	xanthopygius	13	OM	1	1	D
Aves	Passeriformes	Dicruridae	Dicrurus	aeneus	12	IN	1	0	D
Aves	Passeriformes	Dicruridae	Dicrurus	hottentottus	6	IN	1	0	D
Aves	Passeriformes	Dicruridae	Dicrurus	leucophaeus	$\tilde{5}$	IN	1	Õ	D
Aves	Passeriformes	Dicruridae	Dicrurus	paradiseus	18	IN	1	0	D
Aves	Passeriformes	Eurylaimidae	Calyptomena	hosei	4	OM	1	1	D
Aves	Passeriformes	Eurylaimidae	Calyptomena	viridis	16	OM	1	1	D
Aves	Passeriformes	Eurylaimidae	Calyptomena	whiteheadi	4	OM	1	1	D
Aves	Passeriformes	Eurylaimidae	Cymbirhynchus	macrorhynchos	14	OM	1	1	D
Aves	Passeriformes	Eurylaimidae	Eurylaimus	javanicus	15	IN	1	0	D
Aves	Passeriformes	Eurylaimidae	Eurylaimus	ochromalus	18	IN	1	0	D
Aves	Passeriformes	Eurylaimidae	Psarisomus	dalhousiae	2	IN	1	0	D
Aves	Passeriformes	Hirundinidae	Hirundo	tahitica	$14^{-1}$	IN	1	0	D
Aves	Passeriformes	Irenidae	Irena	puella	13	OM	1	1	D
Aves	Passeriformes	Monarchidae	Hypothymis	azurea	16	IN	1	0	D
Aves	Passeriformes	Monarchidae	Terpsiphone	paradisi	18	IN	1	Õ	D
Aves	Passeriformes	Muscicapidae	Copsychus	malabaricus	14	OM	1	1	D
Aves	Passeriformes	Muscicapidae	Copsychus	stricklandii	4	OM	1	1	D
Aves	Passeriformes	Muscicapidae	Culicicapa	ceylonensis	9	IN	1	0	D
Aves	Passeriformes	Muscicapidae	Cyornis	banyumas	5	IN	1	ů 0	D
Aves	Passeriformes	Muscicapidae	Cyornis	caerulatus	6	IN	1	Õ	D
Aves	Passeriformes	Muscicapidae	Cyornis	concretus	8	IN	1	ů 0	D
Aves	Passeriformes	Muscicapidae	Cyornis	rufigastra	3	IN	1	Õ	D
Aves	Passeriformes	Muscicapidae	Cyornis	superbus	13	IN	1	0	D
Aves	Passeriformes	Muscicapidae	Cyornis	turcosus	10	IN	1	0	D
Aves	Passeriformes	Muscicapidae	Cyornis	unicolor	7	IN	1	0	D
Aves	Passeriformes	Muscicapidae	Eumyias	indigo	3	OM	1	1	D
Aves	Passeriformes	Muscicapidae	Eumyias	thalassinus	5	IN	1	0	D
Aves	Passeriformes	Muscicapidae	Ficedula	dumetoria	11	IN	1	0	D
Aves	Passeriformes	Muscicapidae	Ficedula	westermanni	6	IN	1	0	D
Aves	Passeriformes	Muscicapidae	Muscicapella	hodgsoni	4	IN	1	0	D
Aves	Passeriformes	Muscicapidae	Rhinomyias	olivaceus	4	IN IN	1	0	D
Aves	Passeriformes	Muscicapidae	Rhinomyias	ruficauda	47	IN	1	0	D
	Passeriformes			umbratilis		IN IN	1	0	D
Aves	Passeriformes	Muscicapidae	Rhinomyias Trichixos		$\frac{16}{15}$	IN IN		0	D
Aves		Muscicapidae		pyrropygus			1		
Aves	Passeriformes	Nectariniidae	Aethopyga	siparaja	14	OM	1	0	D

Class	Order	Family	Genus	Species	No. of Sites	Guild	Invert	Fruit	Act
Aves	Passeriformes	Nectariniidae	Aethopyga	temminckii	11	ОМ	1	1	D
Aves	Passeriformes	Nectariniidae	Anthreptes	malacensis	10	OM	1	1	D
Aves	Passeriformes	Nectariniidae	Anthreptes	rhodalaemus	8	OM	1	1	D
Aves	Passeriformes	Nectariniidae	Anthreptes	simplex	11	OM	1	1	D
Aves	Passeriformes	Nectariniidae	Arachnothera	affinis	12	OM	1	0	D
Aves	Passeriformes	Nectariniidae	Arachnothera	chrysogenys	12	OM	1	1	D
Aves	Passeriformes	Nectariniidae	Arachnothera	crassirostris	14	OM	1	0	D
Aves	Passeriformes	Nectariniidae	Arachnothera	flavigaster	14	OM	1	1	D
Aves	Passeriformes	Nectariniidae	Arachnothera	juliae	5	NA	0	0	D
Aves	Passeriformes	Nectariniidae	Arachnothera	longirostra	17	OM	1	0	D
Aves	Passeriformes	Nectariniidae	Arachnothera	modesta	1	OM	1	0	D
Aves	Passeriformes	Nectariniidae	Arachnothera	robusta	13	OM	1	0	D
Aves	Passeriformes	Nectariniidae	Chalcoparia	singalensis	13	OM	1	1	D
Aves	Passeriformes	Nectariniidae	Cinnyris	jugularis	8	OM	1	0	D
Aves	Passeriformes	Nectariniidae	Hypogramma	hypogrammica	14	OM	1	1	D
Aves	Passeriformes	Nectariniidae	Leptocoma	calcostetha	6	IN	1	0	D
Aves	Passeriformes	Nectariniidae	Leptocoma	sperata	12	NA	0	0	NA
Aves	Passeriformes	Oriolidae	Oriolus	chinensis	2	OM	1	1	D
Aves	Passeriformes	Oriolidae	Oriolus	cruentus	$\overline{2}$	OM	1	1	D
Aves	Passeriformes	Oriolidae	Oriolus	xanthonotus	$15^{-}$	OM	1	1	D
Aves	Passeriformes	Oriolidae	Oriolus	xanthornus	1	OM	1	1	D
Aves	Passeriformes	Pachycephalidae	Pachycephala	cinerea (grisola)	5	IN	1	0	Ď
Aves	Passeriformes	Pachycephalidae	Pachycephala	hypoxantha	5	IN	1	Ő	D
Aves	Passeriformes	Paridae	Parus	major	1	OM	1	Ő	D
Aves	Passeriformes	Pityriaseidae	Pityriasis	gymnocephala	13	IN	1	ů 0	D
Aves	Passeriformes	Pycnonotidae	Criniger	bres	13	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Criniger	finschii	10	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Criniger	ochraceus	8	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Criniger	phaeocephalus	14	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Hemixos	flavala	8	NA	0	0	D
Aves	Passeriformes	Pycnonotidae	Iole	olivacea	12	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Ixos	malaccensis	$12 \\ 12$	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Pycnonotus	atriceps	15	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Pycnonotus	brunneus	15	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Pycnonotus	cyaniventris	14	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Pycnonotus	e e	14 12	OM	1	1	D
	Passeriformes		v	erythropthalmos eutilotus	$12 \\ 15$	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Pycnonotus Duomonotus		10 3	OM	1	1	D
Aves		Pycnonotidae	Pycnonotus	melanicterus (flaviventris)					
Aves	Passeriformes	Pycnonotidae	Pycnonotus	melanoleucos	9	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Pycnonotus	nieuwenhuisii	3	$\mathbf{FR}$	0	0	D
Aves	Passeriformes	Pycnonotidae	Pycnonotus	plumosus	12	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Pycnonotus	simplex	14	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Pycnonotus	squamatus	8	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Setornis	criniger	9	OM	1	1	D
Aves	Passeriformes	Pycnonotidae	Tricholestes	criniger	13	OM	1	1	D
Aves	Passeriformes	Rhipiduridae	Rhipidura	albicoll is	6	IN	1	0	D
Aves	Passeriformes	Rhipiduridae	Rhipidura	javanica	14	IN	1	0	D
Aves	Passeriformes	Rhipiduridae	Rhipidura	perlata	14	IN	1	0	D
Aves	Passeriformes	Silviidae	Abroscopus	superciliaris	8	IN	1	0	D
Aves	Passeriformes	Silviidae	Phylloscopus	trivirgatus	4	IN	1	0	D
Aves	Passeriformes	Silviidae	Seicercus	montis	4	IN	1	0	D
Aves	Passeriformes	Sittidae	Sitta	frontalis	15	IN	1	0	D
Aves	Passeriformes	Sturnidae	Aplonis	panayensis	7	$\mathbf{FR}$	0	1	D
Aves	Passeriformes	Sturnidae	Gracula	religiosa	16	OM	1	1	D
Aves	Passeriformes	Timaliidae	Alcippe	brunneicauda	15	OM	1	1	D
Aves	Passeriformes	Timaliidae	Garrulax	lugubris (calva)	3	IN	1	0	D

Class	Order	Family	Genus	Species	No. of Sites	Guild	Invert	Fruit	Act
Aves	Passeriformes	Timaliidae	Garrulax	palliatus	5	OM	1	1	D
Aves	Passeriformes	Timaliidae	Macronous	ptilosus	17	IN	1	0	D
Aves	Passeriformes	Timaliidae	Mala copteron	affine	13	IN	1	0	D
Aves	Passeriformes	Timaliidae	Mala copteron	albogulare	9	NA	0	0	D
Aves	Passeriformes	Timaliidae	Mala copteron	cinereum	15	IN	1	0	D
Aves	Passeriformes	Timaliidae	Mala copteron	magnirostre	16	IN	1	0	D
Aves	Passeriformes	Timaliidae	Mala copteron	magnum	16	IN	1	0	D
Aves	Passeriformes	Timaliidae	Pteruthius	flaviscapis	4	OM	1	1	D
Aves	Passeriformes	Timaliidae	Stachyris	ery throp tera	16	IN	1	0	D
Aves	Passeriformes	Timaliidae	Stachyris	leucotis	5	IN	1	0	D
Aves	Passeriformes	Timaliidae	Stachyris	maculata	15	IN	1	0	D
Aves	Passeriformes	Timaliidae	Stachyris	nigriceps	8	IN	1	0	D
Aves	Passeriformes	Timaliidae	Stachyris	nigricollis	14	IN	1	0	D
Aves	Passeriformes	Timaliidae	Stachyris	poliocephala	12	IN	1	0	D
Aves	Passeriformes	Timaliidae	Stachyris	rufifrons	6	IN	1	0	D
Aves	Passeriformes	Timaliidae	Trichastoma	abboti	6	IN	1	0	D
Aves	Passeriformes	Timaliidae	Trichastoma	bicolor	13	IN	1	0	D
Aves	Passeriformes	Timaliidae	Trichastoma	perspicillatum	1	NA	0	0	D
Aves	Passeriformes	Timaliidae	Trichastoma	sepiaria	11	IN	1	0	D
Aves	Passeriformes	Timaliidae	Yuhina	everetti	7	IN	1	0	D
Aves	Passeriformes	Turdidae	Chlamydochaera	jefferyi	4	OM	1	1	D
Aves	Passeriformes	Zosteropidae	Chlorocharis	emiliae	4	OM	1	0	D
Aves	Passeriformes	Zosteropidae	Zosterops	atricapilla	$\overline{2}$	OM	1	1	D
Aves	Passeriformes	Zosteropidae	Zosterops	everetti	3	IN	1	0	D
Aves	Passeriformes	Zosteropidae	Zosterops	flavus	1	IN	1	0 0	D
Aves	Passeriformes	Zosteropidae	Zosterops	palpebrosus	2	IN	1	0	D
Aves	Piciformes	Indicatoridae	Indicator	archipelagicus	6	IN	1	0	D
Aves	Piciformes	Picidae	Blythipicus	rubiginosus	15	IN	1	0	D
Aves	Piciformes	Picidae	Celeus	brachyurus	$13 \\ 12$	OM	1	1	D
Aves	Piciformes	Picidae	Chrysocolaptes	lucidus	12	IN	1	0	D
Aves	Piciformes	Picidae	Dendrocopos	canicapillus	10	OM	1	1	D
	Piciformes	Picidae	Dinopium	javanense	4	IN	1	0	D
Aves Aves	Piciformes	Picidae	Dinopium Dinopium	rafflesii	$12^{4}$	IN IN	1	0	D
	Piciformes	Picidae	*	,,	12	IN IN	1	0	D
Aves	Piciformes		Dryocopus	javensis		OM			D
Aves		Picidae Picidae	Hemicircus Maintanta	concretus	14	IN IN	1	1	D
Aves	Piciformes		Meiglyptes	tristis	16		1	0	
Aves	Piciformes	Picidae	Meiglyptes	tukki	16	IN	1	0	D
Aves	Piciformes	Picidae	Mulleripicus	pulverulentus	15	IN	1	0	D
Aves	Piciformes	Picidae	Picus	mentalis	12	OM	1	1	D
Aves	Piciformes	Picidae	Picus	miniaceus	11	IN	1	0	D
Aves	Piciformes	Picidae	Picus	puniceus	17	IN	1	0	D
Aves	Piciformes	Picidae	Reinwardtipicus	validus	17	IN	1	0	D
Aves	Piciformes	Ramphastidae	Calorhamphus	fuliginosus	15	OM	1	1	D
Aves	Piciformes	Ramphastidae	Megalaima	australis	16	OM	1	1	D
Aves	Piciformes	Ramphastidae	Megalaima	chry sopogon	12	OM	1	1	D
Aves	Piciformes	Ramphastidae	Megalaima	eximia	8	$\mathbf{FR}$	0	1	D
Aves	Piciformes	Ramphastidae	Megalaima	henricii	12	$\mathbf{FR}$	0	1	D
Aves	Piciformes	Ramphastidae	Megalaima	monticola	7	OM	1	1	D
Aves	Piciformes	Ramphastidae	Megalaima	mystacophanos	16	OM	1	1	D
Aves	Piciformes	Ramphastidae	Megalaima	pulcherrima	4	OM	1	1	D
Aves	Piciformes	Ramphastidae	Megalaima	rafflesii	14	OM	1	1	D
Aves	Psittaciformes	Psittacidae	Loriculus	galgulus	16	$\mathbf{FR}$	0	1	D
Aves	Psittaciformes	Psittacidae	Psittacula	longicauda	8	$\mathbf{FR}$	0	1	D
Aves	Psittaciformes	Psittacidae	Psittinus	cyanurus	10	$\mathbf{FR}$	0	1	D
Aves	Psittaciformes	Psittacidae	Tanygnathus	lucionensis	2	$\mathbf{FR}$	0	1	D
Aves	Strigiformes	Strigidae	Otus	rufenscens	8	IN	1	0	Ν
Aves	Trogoniformes	Trogonidae	Harpactes	dardii	14	IN	1	0	D
Aves	Trogoniformes	Trogonidae	Harpactes	duvaucelii	16	NA	0	0	D
	Trogoniformes	Trogonidae	Harpactes	kasumba	14	OM	1	1	D

Class	Order	Family	Genus	Species	No. of Sites	Guild	Invert	Fruit	Ac
Aves	Trogoniformes	Trogonidae	Harpactes	oreskios	8	NA	0	0	D
Aves	Trogoniformes	Trogonidae	Harpactes	orrophaeus	9	NA	0	0	D
Aves	Trogoniformes	Trogonidae	Harpactes	whiteheadi	<b>5</b>	IN	1	0	D
Aves	(unclear)	INCERTAE SEDIS	Erporn is	zantholeuca	10	OM	1	1	D
Aves	(unclear)	INCERTAE SEDIS	Orthotomus	atrogularis	14	IN	1	0	D
Aves	(unclear)	INCERTAE SEDIS	Orthotomus	ruficeps	13	IN	1	0	D
Aves	(unclear)	INCERTAE SEDIS	Philentoma	pyrrhoptera	15	IN	1	0	D
Aves	(unclear)	INCERTAE SEDIS	Philentoma	velata (velatum)	11	IN	1	0	D
Aves	(unclear)	INCERTAE SEDIS	Tephrodorn is	gularis	14	IN	1	0	D
Aves	(unclear)	Podargidae	Batrachostomus	cornutus	3	NA	0	0	Ν
Mammalia	Chiroptera	Emballonuridae	Emballonura	a lecto	10	NA	0	0	Ν
Mammalia	Chiroptera	Emballonuridae	Emballonura	monticola	9	NA	0	0	Ν
	Chiroptera	Hipposideridae	Coelops	robinsoni	$^{2}$	IN	1	0	Ν
	Chiroptera	Hipposideridae	Hipposideros	ater	6	IN	1	0	Ν
	Chiroptera	Hipposideridae	Hipposideros	bicolor	4	IN	1	0	Ν
	Chiroptera	Hipposideridae	Hipposideros	cervinus	11	IN	1	0	Ν
	Chiroptera	Hipposideridae	Hipposideros	cineraceus	6	IN	1	0	Ν
	Chiroptera	Hipposideridae	Hipposideros	diadema	8	IN	1	0	N
	Chiroptera	Hipposideridae	Hipposideros	doriae (sabanus)	3	IN	1	Õ	N
	Chiroptera	Hipposideridae	Hipposideros	dyacorum	6	IN	1	ů 0	N
	Chiroptera	Hipposideridae	Hipposideros	galeritus	9	IN	1	0	N
	Chiroptera	Hipposideridae	Hipposideros	larvatus	4	IN	1	0	N
	Chiroptera	Hipposideridae	Hipposideros	ridleyi	5	IN	1	0	N
	Chiroptera	Nycteridae	Nycteris	tragata (javanica)	8	IN	1	0	N
	Chiroptera	Pteropodidae	Aethalops	aequalis (alecto)	4	FR	0	1	N
	Chiroptera	Pteropodidae	Balionycteris	maculata	$15^{4}$	FR	0	1	N
	Chiroptera	Pteropodidae	Chironax		3	FR	0	1	N
				melanocephalus		гл FR		1	N
	Chiroptera	Pteropodidae	Cynopterus	brachyotis	15		0		
	Chiroptera	Pteropodidae	Cynopterus	horsfieldii	3	FR	0	1	N
	Chiroptera	Pteropodidae	Cynopterus	sphinx	1	NA	0	0	N
	Chiroptera	Pteropodidae	Dyacopterus	spadiceus	5	FR	0	1	N
	Chiroptera	Pteropodidae	Eonycteris	major	1	FR	0	1	N
	Chiroptera	Pteropodidae	Eonycteris	spelaea	7	FR	0	1	N
	Chiroptera	Pteropodidae	Macroglossus	minimus	11	FR	0	1	N
	Chiroptera	Pteropodidae	Megaerops	ecaudatus	5	FR	0	1	N
	Chiroptera	Pteropodidae	Megaerops	wetmorei	2	FR	0	1	N
	Chiroptera	Pteropodidae	Penthetor	lucasi	8	FR	0	1	Ν
	Chiroptera	Pteropodidae	Pteropus	vampyrus	13	$\mathbf{FR}$	0	1	Ν
	Chiroptera	Pteropodidae	Rousettus	amplexicaudatus	2	$\mathbf{FR}$	0	1	Ν
	Chiroptera	Pteropodidae	Rousettus	spinalatus	1	$\mathbf{FR}$	0	1	Ν
	Chiroptera	Rhinolophidae	Rhinolophus	a cuminatus	3	IN	1	0	Ν
	Chiroptera	Rhinolophidae	Rhinolophus	affinis	4	IN	1	0	Ν
	Chiroptera	Rhinolophidae	Rhinolophus	borneensis	15	IN	1	0	Ν
	Chiroptera	Rhinolophidae	Rhinolophus	creaghi	4	IN	1	0	Ν
Mammalia	Chiroptera	Rhinolophidae	Rhinolophus	luctus	7	IN	1	0	Ν
	Chiroptera	Rhinolophidae	Rhinolophus	philippinensis	$^{2}$	IN	1	0	Ν
Mammalia	Chiroptera	Rhinolophidae	Rhinolophus	pusillus	$^{2}$	IN	1	0	Ν
Mammalia	Chiroptera	Rhinolophidae	Rhinolophus	sedulus	12	IN	1	0	Ν
Mammalia	Chiroptera	Rhinolophidae	Rhinolophus	trifoliatus	14	IN	1	0	Ν
Mammalia	Chiroptera	Vespertilionidae (Kerivoulininae)	Kerivoula	hardwickii	8	IN	1	0	Ν
	Chiroptera	Vespertilionidae (Kerivoulininae)	Kerivoula	intermedia	9	IN	1	0	N
Mammalia	Chiroptera	Vespertilionidae (Kerivoulininae)	Kerivoula	minuta	7	IN	1	0	Ν
Mammalia	Chiroptera	Vespertilionidae (Kerivoulininae)	Kerivoula	papillosa	12	IN	1	0	Ν
Mammalia	Chiroptera	Vespertilionidae (Kerivoulininae)	Kerivoula	pellucida	8	IN	1	0	Ν

Class	Order	Family	Genus	Species	No. of Sites	Guild	Invert	Fruit	Act
Mammalia	Chiroptera	Vespertilionidae (Kerivoulininae)	Kerivoula	whiteheadii	1	IN	1	0	N
Mammalia	Chiroptera	(Kerivoulininae) (Kerivoulininae)	Phoniscus	atrox	4	IN	1	0	Ν
Mammalia	Chiroptera		Phoniscus	jagorii	2	IN	1	0	Ν
Mammalia	Chiroptera	Vespertilionidae (Murininae)	Harpiocephalus	harpia	1	IN	1	0	Ν
Mammalia	Chiroptera	Vespertilionidae (Murininae)	Murina	aenea	3	IN	1	0	Ν
	Chiroptera	(Murininae)	Murina	cyclotis	4	IN	1	0	Ν
Mammalia	-	Vespertilionidae (Murininae)	Murina	rozendaali	3	IN	1	0	Ν
Mammalia	Chiroptera	Vespertilionidae (Murininae)	Murina	suilla	10	IN	1	0	Ν
Mammalia	Primates	Cercopithecidae	Macaca	fascicularis	21	OM	1	1	D
Mammalia	Primates	Cercopithecidae	Nasalis	larvatus	10	FO	0	1	D
Mammalia	Primates	Cercopithecidae	Presbytis	cristata	10	FO	0	1	D
Mammalia	Primates	Cercopithecidae	Presbytis	frontata	6	FO	0	0	D
Mammalia	Primates	Cercopithecidae	Presbytis	hosei	7	FO	0	0	D
Mammalia	Primates	Cercopithecidae	Presbytis	melalophos	3	FO	0	0	D
Mammalia	Primates	Cercopithecidae	Presbytis	rubicunda	16	FO	0	0	D
Mammalia	Primates	Hominidae	Pongo	pygmaeus	13	$\mathbf{FR}$	1	1	D
Mammalia	Primates	Hylobatidae	Hylobates	agilis	5	FRFO	0	1	D
Mammalia	Primates	Hylobatidae	Hylobates	muelleri	17	$\mathbf{FR}$	0	1	D
Mammalia	Primates	Lorisidae	Nycticebus	coucang	12	OM	1	1	Ν
Mammalia	Primates	Tarsiidae	Tarsius	bancanus	14	IN	1	0	Ν
Mammalia	Rodentia	Sciuridae	Aeromys	tephromelas	4	FRFO	0	1	Ν
Mammalia	Rodentia	Sciuridae	Aeromys	thomasi	4	$\mathbf{FR}$	0	1	Ν
Mammalia	Rodentia	Sciuridae	Callosciurus	adamsi	3	NA	0	0	D
Mammalia	Rodentia	Sciuridae	Callosciurus	baluensis	3	NA	0	0	D
Mammalia	Rodentia	Sciuridae	Callosciurus	notatus	16	OM	1	1	D
Mammalia	Rodentia	Sciuridae	Callosciurus	orestes (formerly nigrovittatus)	4	OM	1	1	D
Mammalia	Rodentia	Sciuridae	Callosciurus	prevostii	19	OM	1	1	D
Mammalia	Rodentia	Sciuridae	Exilisciurus	exilis	15	OM	1	0	D
Mammalia	Rodentia	Sciuridae	Exilisciurus	whiteheadi	6	FO	0	0	NA
Mammalia	Rodentia	Sciuridae	Glyphotes	simus	3	NA	0	0	D
Mammalia	Rodentia	Sciuridae	Hylopetes	lepidus	1	NA	0	0	N
Mammalia	Rodentia	Sciuridae	Hylopetes	spadiceus	1	NA	0	0	N
Mammalia	Rodentia	Sciuridae	Iomys	horsfieldi	3	NA	0	0	N
Mammalia	Rodentia	Sciuridae	Nannosciurus	melanotis	12	NA	0	0	D
Mammalia	Rodentia	Sciuridae	Petaurista	elegans	1	NA	0	0	N
Mammalia	Rodentia	Sciuridae	Petaurista	petaurista	5	FRFO	0	1	N
Mammalia	Rodentia	Sciuridae	Petinomys	genibarbis	3	NA	0	0	N
Mammalia	Rodentia	Sciuridae	Petinomys	hageni	1	NA	0	0	N
Mammalia	Rodentia Rodentia	Sciuridae Sciuridae	Petinomys Patinomuo	setosus	1	NA NA	0	0	N N
Mammalia Mammalia	Rodentia Rodentia	Sciuridae Sciuridae	Petinomys Pteromysous	vordermanni	1	NA NA	0	0	N N
Mammalia	Rodentia	Sciuridae	Pteromyscus Patufa	pulverulentus affinio	1 18	NA FRFO	0 0	0	N D
Mammalia Mammalia	Rodentia	Sciuridae Sciuridae	Ratufa Sundasciurus	affinis brookei	18 6	NA	0	0 0	D
Mammalia	Rodentia	Sciuridae	Sundasciurus Sundasciurus		6 14	OM	1	1	D
Mammalia	Rodentia	Sciuridae	Sundasciurus Sundasciurus	hippurus jentinki	14 4	NA	0	0	D
Mammalia	Rodentia	Sciuridae	Sundasciurus Sundasciurus	lowii	$\frac{4}{14}$	OM	1	1	D
Mammalia		Sciuridae	Sundasciurus Sundasciurus	tenuis	$\frac{14}{7}$	OM	1	1	D

Species attribute data and taxonomy: guild assignments include insectivore (IN), omnivore (OM), frugivore (FR), folivore (FO), frugivore/folivore (FRFO), and unknown (NA). Invert and fruit columns indicate the presence (1) or absence (0) of invertebrates or fruits in the diet respectively. Activity patterns include diurnal (D), nocturnal (N), or unknown (NA).

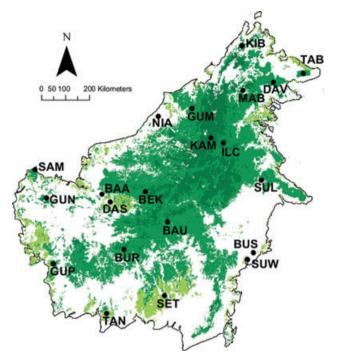


Fig. 2. Locations of sites on Borneo in Southeast Asia: BAU: Barito Ulu; BAA: Batang Ai National Park; BEK\*: Betung Kerihun National Park; BUR\*: Bukit Raya-Bukit Baka National Park; BUS: Bukit Soeharto Wildlife Reserve; DAS: Danau Sentarum National Park; DAV: Danum Valley Conservation Forest; GUM: Gunung Mulu National Park; GUN: Gunung Niut Wildlife Reserve; GUP: Gunung Palung National Park; ILC: Inhutani Logging Concession; KAM\*: Kayan Mentarang National Park; KIB\*: Kinabalu National Park; MAB\*: Maliau Basin Conservation Area; NIA: Niah National Park; SAM: Samunsam Wildlife Sanctuary; SET: Sebangau National Park; SUL\*: Sungai Lesan Protection Forest; SUW: Sungai Wain Protection Forest; TAB: Tabin Wildlife Reserve; TAN: Tanjung Puting National Park. Sites with similar forest types included in the habitat analysis are indicated with an asterisk. Forest cover according to Miettinen et al. [2011]. Light shading represents swamp forests and mangroves, dark shading represents montane forest and all other shading represents lowland forests.

matter, in which case we classified the species as an omnivore.

All data compilation and analysis for this study took place at the University of California, Davis beginning in August 2010. We collected no new field data and as such there were no applicable institutional animal care committee protocols to be approved. All research adhered to the legal requirements of the United States and to the American Society of Primatologists principles for the ethical treatment of nonhuman primates.

#### Analysis

Our objective was to determine the extent of ecological structure present in communities of primates and in the broader vertebrate communities in which primates live. We therefore investigated each co-occurrence pattern using all species combined as well as several subsets of the data, which included analyzing each taxonomic group individually (primates, birds, bats, or squirrels), or grouped as nonvolant species (primates and squirrels), mammals (primates, squirrels and bats), diurnal species (diurnal primates, squirrels and birds), and all species.

There has been substantial debate on the subject of determining  $\alpha$  values in statistical testing [Rice, 1989; Roback & Askins, 2005]. We agree with the argument that authors should reject correction factors in ecological studies and instead report exact *P*-values and interpret results reasonably [Moran, 2003; Nakagawa, 2004]. We therefore report exact *P*-values and base our interpretation of the data on the  $\alpha = 0.05$  level.

## **Checkerboard Distributions**

To test for checkerboard distributions, we used the co-occurrence analysis in the program EcoSim [Gotelli & Entsminger, 2011] to calculate C-scores [Stone & Roberts, 1990]. We compared observed scores to a null distribution model based on 5,000 permutations with fixed column and row totals. A C-score measures the mean number of "checkerboard units" based on all possible pairs of species in a presence-absence matrix. A checkerboard unit refers to a  $2 \times 2$  site by species submatrix in which one site contains only one species and the other site only contains the other species such that the two species do not co-occur at either site [Gotelli & Entsminger, 2011]. EcoSim follows Stone and Roberts [1990] in calculating a checkerboard unit as  $CU_{ij} = (\mathbf{r}_i - S_{ij})(\mathbf{r}_j - S_{ij})$ , where  $\mathbf{r}_i$  and  $\mathbf{r}_j$  are the total number of occurrences across sites of species *i* and *j* and S is the number of sites where the two species co-occur. The *C*-score is the mean of the *CU*-values for all species pairs in a matrix. If the observed matrix has a significantly higher C-score than matrices under the null model, this suggests evidence of significant ecological structure in the community, which can result from species segregation, aggregation, or turnover [Ulrich and Gotelli, 2012]. While even the most conservative measure suggests that significant C-scores are typically the result of species segregation, which is consistent with the traditional C-score interpretation of competitive exclusion, further analvses at the species-pair level are necessary to determine the causal factor [Ulrich and Gotelli, 2012]. If nonprimate vertebrate species importantly interact with primate species, then we expect to find higher C-scores when we include nonprimates in the analysis than when we examine primates alone.

Because species responses to variation in habitats (i.e. habitat checkerboards) can also cause significant *C*-scores [Gotelli & McCabe, 2002], we performed additional analyses to control for the effects of variation in habitat on checkerboard distributions. We conducted the analysis with a subset of six of the total sites because they encompassed the same

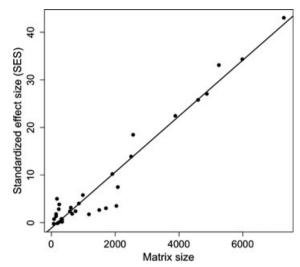


Fig. 3. Linear relationship between matrix size and SES in C-score analysis for all communities. There is a significant positive linear relationship between the SES and the size of the presence–absence matrix, which is defined as the number of species times the number of sites (linear regression, df = 30,  $R^2 = 0.94$ , P < 0.001). Because the SES scales with matrix size, we limit direct comparisons of SES values and instead rely on the patterning of significant C-score values across communities.

broad habitat types of lowland, hill, and montane forest. To examine the effects of interactions related to specific types of resources, we also conducted the co-occurrence analysis separately using only species whose diets include invertebrates and again using only species whose diets include fruit.

For all checkerboard analyses, we report the standardized effect size (SES) and the probability that the observed C-score is greater than the expected C-score given the simulated data (P). We calculate the SES as the (observed index - mean of simulated indices) / standard deviation of simulated indices. Published guidelines note that the SES allows for comparisons between different tests because it scales the results of each test in units of standard deviations [Gotelli & Entsminger, 2011]. However, there is a significant positive linear relationship between the SES and the size of the presence-absence matrix, which is defined as the number of species times the number of sites (linear regression, df =30,  $R^2 = 0.94$ , P < 0.001; Fig. 3). We therefore do not interpret relative differences in SES values between communities as biologically meaningful. We focus on the comparison of expected and observed *C*-score values for each community and we examine the patterning of significantly checkerboarded communities based on analyses of all sites and when we controlled for habitat and diet characteristics.

## **Guild Proportionality**

To test for guild proportionality, we conducted G-tests of independence [McDonald, 2009] using the function g.test [Hurd, 2011] in the software R 2.13.0

TABLE II. Number of species included in analyses

	Bats	Birds	Primates	Squirrels
(a) Foraging guild				
Folivores	0	0	6	1
Frugivores	15	23	2	1
Frugivore/Folivores	0	0	1	3
Insectivores	34	112	1	0
Omnivores	0	94	2	7
Unknown	3	11	0	15
Total	52	240	12	27
(b) Diet				
Fruit	15	103	7	9
Invertebrates	34	205	4	7

Number of species included in analyses for each taxonomic group (a) for each foraging guild used in the analyses for guild proportionality and Fox's assembly rule for favored states and (b) with diets containing fruits or invertebrates used in the analyses for checkerboard distributions based on species whose diets contained fruit or invertebrates, respectively. Overall checkerboard analyses included species from all foraging guilds including unknown, but guild proportionality analyses excluded species with unknown foraging guilds.

[R Development Core Team, 2012]. If the proportion of species in guilds changes significantly across sites, then P < 0.05. If the proportion of species in guilds is fixed, then P > 0.95. If 0.05 < P < 0.95, then the proportion of guilds remains constant but with noise [Feeley, 2003]. If nonprimate vertebrates are more important competitors of primates than other primates, then we expect to find greater evidence of guild proportionality when we include nonprimate vertebrates. We excluded species for which dietary data were unavailable from the guild-level analyses (Table II).

#### Fox's Assembly Rule

We compared the observed number of favored states to the distribution of favored states expected under the null model using the favored states option within the guild analysis in EcoSim [Gotelli & Entsminger, 2011]. We implemented the recommended program settings for column and row totals. If nonprimate vertebrates are stronger competitors of primates than other primates, then we expect to find greater evidence in support of Fox's assembly rule when we consider nonprimate vertebrates at the same time as primates than when we consider primates in isolation.

#### RESULTS

#### **Checkerboard Distributions**

Significantly high checkerboard scores are consistent with the interpretation that species interactions have structured co-occurrence patterns across communities. When we included all sites (N = 21), observed checkerboard scores were greater than expected for all communities analyzed (Fig. 4); this

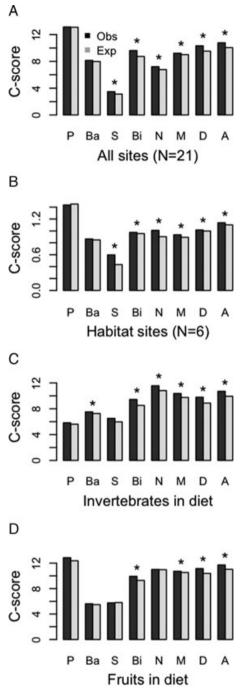


Fig. 4. Comparison of observed and expected checkerboard scores in dark gray and light gray, respectively, for (A) all sites (B) and sites with similar habitat (C) and for fruit-eating species (B) and (D) invertebrate eating species (Ba: bats; Bi: birds; P: primates; S: squirrels; N: nonvolant; M: mammals; D: diurnal; A: all species). If an observed matrix has a significantly higher C-score than expected given the null model indicated by an asterisk, then species pairs co-occur less often than expected by chance. When interpreting the graph of the observed versus expected c-scores, it is the relative difference between observed and the expected rather than the absolute c-score values that is meaningful. Significantly high C-scores are consistent with the interpretation that there is significant ecological structure within communities.

patterning was significant when we analyzed individual taxonomic groups of birds (SES = 33.10, P < 0.001) or squirrels (SES = 3.15, P < 0.001), but not bats (SES = 1.74, P = 0.053) or primates (SES = 0.14, P = 0.437). In the communities with taxonomic groups combined, checkerboard scores were significant for all combinations. This included significant *C*-scores for nonvolant species (SES = 4.01, P = 0.002), mammals (SES = 3.50, P < 0.001), diurnal species (SES = 34.32, P < 0.001), and all species combined (SES = 43.04, P < 0.001; Fig. 4A).

In the analysis of sites with similar habitat (N = 6), the patterns were consistent with the results for all sites. Observed checkerboard scores were greater than those expected under the null models for all communities analyzed except for primates; this patterning was significant when we analyzed individual taxonomic groups of birds (SES = 2.64, P = 0.017) or squirrels (SES = 5.00, P < 0.001), but not bats (SES = 0.65, P = 0.235) or primates (SES = -0.24, P = 0.604). Among combined taxonomic groups, checkerboard scores were significant for all combinations, including nonvolant species (SES < 3.82, P = 0.003), mammals (SES = 2.38, P = 0.025), diurnal species (SES = 3.02, P = 0.001) and all species (SES = 7.47, P < 0.001) (Fig. 4B).

We also ran the checkerboard analysis for a subset of species based on the presence of broad dietary items in their diets. For species with fruit in their diets, birds were the only taxonomic group with significant checkerboard distributions when considered in isolation (SES = 10.21, P < 0.001). Although primates considered in isolation were marginally significant (SES = 1.79, P = 0.060), the other taxonomic groups did not exhibit significant checkerboard scores (bats: SES = 0.78, P = 0.222); squirrels: SES = -0.15, P = 0.522) nor did the nonvolant species (SES = 0.16, P = 0.400). We found significantly high checkerboard scores for fruit-eating mammals (SES = 1.84, P = 0.046), diurnal species (SES = 13.88, P < 0.001), and all species combined (SES = 18.46, P < 0.001; Fig. 4C).

For species with invertebrates in their diets, checkerboard distributions were significant in communities of bats (SES = 2.39, P = 0.015), birds (SES = 22.43, P < 0.001), mammals (SES = 5.78, P < 0.001), nonvolant species (SES = 2.82, P = 0.012), diurnal species (SES = 25.79, P < 0.001), and all species combined (SES = 27.05, P < 0.001), but not in primates (SES = 0.73, P < 0.241) or squirrels (SES = 1.35, P = 0.110) (Fig. 4D).

## **Guild Proportionality**

We found results consistent with "fixed" guild proportionality in primates (G = 32.70,  $\chi^2 df = 80$ , P = 1.00), squirrels (G = 28.02,  $\chi^2 df = 60$ , P = 1.00), birds (G = 19.14,  $\chi^2 df = 40$ , P = 0.998), and nonvolant species (primates and squirrels combined;

**TABLE III.** Guild proportionality results

Taxa	Log-likelihood ratio statistic (G)	$\chi^2~degrees$ of freedom	P-value
All species	89.04	80	0.23
Bats	44.06	20	< 0.01
Birds	19.14	40	1.00
Diurnal	66.09	80	0.87
Mammals	108.59	80	0.02
Nonvolant	35.58	80	1.00
Primates	32.70	80	1.00
Squirrels	28.02	60	1.00

Guild proportionality results for *G*-tests of independence. Guild proportionality is consistent with competition at the guild level structuring communities when resources are distributed homogeneously across sites. P < 0.05 suggests the proportion of guilds changes significantly across sites. P > 0.95 suggests the proportion of guilds changes is fixed. If 0.05 < P < 0.95, then the proportion of guilds remains constant but with noise [Feeley, 2003].

G = 35.58,  $\chi^2 df = 80$ , P = 1.00). Significant departure from guild proportionality was evident for bats (G = 44.06,  $\chi^2 df = 20$ , P = 0.001) and mammals (G = 108.59,  $\chi^2 df = 80$ , P = 0.018). Results for guild proportionality in diurnal species (G = 66.09,  $\chi^2 df = 80$ , P = 0.868) and all species combined (G = 89.04,  $\chi^2 df = 80$ , P = 0.229) were nonsignificant (Table III).

## Fox's Assembly Rule for Favored States

We did not find support for Fox's assembly rule for any of the communities (all species, P = 1.00; bats, P = 0.796; birds, P = 0.806; diurnal species, P = 1.00; mammals, P = 0.995). Initial analysis was not applicable for primates, squirrels, or nonvolant vertebrates because these communities did not contain multiple species from each guild. We did not find support for the rule when the guilds with one species or less were omitted (primates, P = 0.989; squirrels, P = 0.222; nonvolant species, P = 0.340).

## DISCUSSION

We evaluated the extent to which species interactions shape vertebrate communities, specifically communities of primates, birds, bats, and squirrels in Borneo. We investigated co-occurrence patterns consistent with interactions structuring the presence or absence of species in communities. Our results suggest that consideration of the broader mammalian and avian communities in which primates live is of central importance for understanding the distribution of primate species and the composition of communities. We found strong support for checkerboard distributions (i.e. the alternating presence or absence of species across sites), particularly when taxonomic groups were combined, and we found the same patterns after accounting for habitat heterogeneity. We also found significant evidence of checkerboard distributions when we analyzed communities defined by dietary characteristics by examining species whose diets include fruits or invertebrates. We found evidence of guild proportionality in birds, primates, squirrels, and the nonvolant species, but did not find support for Fox's assembly rule in any of the communities examined. These results demonstrate the presence of significant ecological structure, which indicates that these communities are nonrandom assemblages of species, and are consistent with the interpretation that species interactions within and between these taxonomic groups may have shaped both species distributions and community structure. Moreover, these results suggest that interactions between primates and other taxonomic groups can have important effects on primate community composition and future research should take account of them.

Variation in the size and habitat heterogeneity of the protected areas is a potential alternative explanation for the significant checkerboard scores because habitat checkerboards, in addition to species interactions, are known to drive checkerboard species patterns [Gotelli et al., 2010; Gotelli & McCabe, 2002]. Given that the sites included in this study cover all major forest types found on Borneo, it is possible that our checkerboard results may reflect habitat checkerboards. Complementary analysis with a subset of the data restricted to similar habitat types, however, yielded consistent results. Thus, while it is likely that habitat heterogeneity contributes to the checkerboard distributions we found when we considered all sites, it is unlikely that it can explain them entirely.

With the notable exceptions of primate communities and bat communities, we found that all other communities examined were significantly checkerboarded when we included species with all diet types. However, we note that communities of insecteating bats were significantly checkerboarded. Previous research on rhinolophoid bats in southeast Asia demonstrated that competition has led to niche differentiation within this insectivorous guild [Kingston et al., 2000], although recent research on North Borneo and the Malay peninsula suggests that insectivorous bat assemblages are relatively homogenous over substantial geographic distances [Struebig et al., In review; Struebig et al., 2011]. Thus, primates are particularly noteworthy for their lack of significant checkerboard distributions when considered in isolation and for the presence of significant C-scores when analyzed with other taxa. These results suggest that there is less interspecific interaction among primate species than between primates and other taxa in Borneo. Primates may be interacting more with nonprimates because of the low diversity of primates in comparison to other taxa.

Past competition may have resulted in sufficient niche differentiation among primates to enable coexistence whereas constraints on competition between primates and other taxa may have prevented further differentiation and thus led to significant checkerboard patterning. Alternatively, historical changes in habitat cover may have driven surviving species into previously unoccupied niches [*sensu* Crowley et al., 2012].

For invertebrate-eating species, checkerboard results were significant for all communities except for primates and squirrels. However, for fruit-eating species, checkerboard scores were not significant for primates, squirrels, bats, or nonvolant species. These results are consistent with the interpretation that invertebrate eating species show more evidence of checkerboard distributions than fruit eating species. In Bornean forests, interactions by species feeding on invertebrates may result in significant ecological patterning more often than interactions for fruit, particularly within taxonomic groups. Because of the extreme fluctuations in the availability of fruit in Bornean forests [Cannon et al., 2007], fruit-eating species may necessarily rely on other resources when fruit is scarce (e.g. orangutans [Knott, 1998]). Invertebrates may represent an important source of fallback foods, sensu Marshall and Wrangham [2007], and consequently function as limiting resources that negatively affect species coexistence. While our study does not address the specific invertebrate and fruit resources that species may be competing for, our results suggest that further investigation of overlap in fallback foods among Bornean vertebrates is warranted.

A handful of recent papers have investigated checkerboard distributions in communities relevant to Bornean vertebrates. A study of primate communities in the Sunda Shelf area found significantly high C-scores for observed primate communities when defined by all species or by folivorous species, but not by frugivorous species [Kamilar & Ledogar, 2011]. The authors suggested that the lack of significant difference in C-scores among frugivorous communities was a reflection of the generalist habits of these species [Kamilar & Ledogar, 2011]. Another recent study investigated checkerboard distributions of mammals on 240 small islands in Southeast Asia [Cardillo & Meijaard, 2010]. When all islands were considered, Cercopithecidae, Viverridae (civets), Pteropodidae (fruit bats), and Sciuridae (squirrels) had significantly higher C-scores than expected under the null model, but Cervidae (deer), Felidae (cats), and Emballonuridae (sheath-tailed bats) did not. While these studies have demonstrated significant co-occurrence patterns within individual families of Southeast Asian mammals, we build upon this growing body of research by considering competitive interactions among families and between mammals and birds.

Our study did not address the additional alternative hypothesis that historical environmental conditions may have caused current checkerboard distributions. In addition, the analyses undertaken in this study assume that species dispersal among sites is not limited. We suggest that future studies explore the role of historical habitats and dispersal ability in driving community composition. Lastly, we suggest that future work investigate potentially artificial effects of sampling effort and seasonality on co-occurrence patterns.

We found fixed guild structure in birds, primates, squirrels, and the nonvolant communities, which supports the test for guild proportionality in these communities. These results are consistent with the interpretation that competition within and between these taxonomic groups has resulted in proportional membership in foraging guilds across sites for squirrels, primates, and the two taxa combined, which differs from the results of the checkerboard analyses and highlights the potential influence of guild characterizations for driving patterns. We also note that guilds containing few species, as is the case with primates and squirrels, may be more likely to appear fixed due to the nature of the statistical test.

Although there is evidence in favor of Fox's assembly rule for communities of invertebrates [Rodriguez-Fernandez et al., 2006], salamanders [Adams, 2007], birds [de Silva & Medellin, 2002], small mammal communities in multiple regions [Fox & Kirkland, 1992; Kelt, 1999; Kelt et al., 1995; McCay et al., 2004], and in Malagasy lemur communities [Ganzhorn, 1997], we did not find support for the rule in any of the vertebrate communities examined. Fox's assembly rule assumes that the distribution of resources is homogeneous within and between sites and it is likely that this assumption is not met given that fruit production varies across habitat types [Marshall et al., 2009a; Wich et al., 2011]. An alternative explanation is that our foraging guild categorization was insufficient to detect competition because a lack of resolution in the dietary data prevented us from a more nuanced analysis. Lastly, we may have failed to find support for Fox's assembly rule because we lacked foraging guild data for some species. The absence of a subset of species may have been sufficient to preclude support of this very strictly defined rule.

Despite the overwhelming complexity and diversity of tropical rain forests, we found evidence of significant ecological structure in Bornean vertebrate communities of primates, birds, bats, and squirrels. This research addresses a gap in our understanding of primate communities; our results contribute to the understanding of the extent to which deterministic interactions within and between taxonomic groups may structure these communities. We suggest that as the field of primate community ecology progresses, it is important that primatologists take into account the broader vertebrate communities in which primates occur and consider the role of biotic interactions within and between taxonomic groups in shaping primate communities. We have drawn attention to the importance of nonprimates as potential competitors of primates with this study, and we hope that more researchers will explicitly consider interspecific interactions between primates and other taxa. We believe that this is an important area of future research that warrants consideration across regions.

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