

# Endemism and diversity of small mammals along two neighboring Bornean mountains

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# **Abstract**

Mountains offer replicated units with large biotic and abiotic gradients in a reduced spatial scale. This transforms them into well-suited scenarios to evaluate biogeographic theories. Mountain biogeography is a hot topic of research and many theories have been proposed to describe the changes in biodiversity with elevation. Geometric constraints, which predict the highest diversity to occur in mid-elevations, have been a focal part of this discussion. Despite this, there is no general theory to explain these patterns, probably because of the interaction among different predictors with the local effects of historical factors. We characterize the diversity of small non-volant mammals across the elevational gradient on Mount (Mt.) Kinabalu (4,095 m) and Mt. Tambuyukon (2,579 m), two neighboring mountains in Borneo, Malaysia. We documented a decrease in species richness with elevation which deviates from expectations of the geometric constraints and suggests that spatial factors (eg. larger diversity in larger areas) are important. The lowland small mammal community was replaced in higher elevations (from above ~1900 m) with montane communities consisting mainly of high elevation Borneo endemics. The positive correlation we find between elevation and endemism is concordant with a hypothesis that predicts higher endemism with topographical isolation. This supports lineage history and geographic history could be important drivers of species diversity in this region.

**Keywords:** Mt. Kinabalu, Mt. Tambuyukon, Shannon index, Sundaland, elevational gradient

# Introduction

Understanding the mechanisms that are responsible for shaping patterns of biodiversity across geography has been an important driver of biological research (eg. Wallace, 1869; Heaney, 1986; Rosenzweig, 1995; Lomolino et al., 2010). Mountains are valuable natural experiments that allow researchers to test biogeographical hypotheses for these reasons: (1) they have limited confounding variation across historical and ecological conditions, (2) they are discrete units to study and (3) they offer replicated gradients for factors (climatic, spatial, ecological) that have been central for research in biogeography (Brown, 2001). Numerous studies have sought to explain the change of alpha diversity across elevation for diverse taxonomic groups (eg. Rahbek, 1995; Patterson & Stotz, 1998; Heaney, 2001; Chen et al., 2006; Kluge, Kessler & Dunn, 2006). Previous research on diversity gradients on mountains has focused on the relationship between diversity gradients on mountains with temperature and precipitation, primary productivity, area, isolation, and geometric constraints. However, the effects of abiotic predictors on mountains are inconsistent across studies (eg. Rahbek, 1995; Patterson & Stotz, 1998; Heaney, 2001; Chen et al., 2006; Kluge, Kessler & Dunn, 2006), and to date, no uniform theory explains mammalian diversity gradients on mountains (Brown, 2001; Lomolino, 2001; Heaney, 2001; Stevens, Rowe & Badgley, 2019).

Geometric constraints stand out from the former predictors as the most recurrent explanation for diversity gradients on mountains across many systems (Rahbek, 1995; Fu et al., 2006; Kluge, Kessler & Dunn, 2006; Rowe, 2009). Geometric constraints predict that the overlap of species' ranges on a constrained area causes species richness to be higher at the center of this area (Colwell & Lees, 2000). Translated to mountains, geometric constraints predict a higher diversity at mid-elevations (mid-elevation bulge) caused by the overlap of species ranges with

midpoints at different elevations. This phenomenon is often referred to as Mid-Domain Effect (MDE). In mammals, a MDE has been reported across different mountain ranges, but it is not a global pattern (Rickart, 2001; Heaney, 2001; Nor, 2001; Li, Song & Zeng, 2003; McCain, 2004, 2005; Rowe, 2009; Rowe, Heaney & Rickart, 2015; Hu et al., 2017).

The history of lineage and place can profoundly influence current patterns of diversity on mountains (Brown, 2001; Lomolino, 2001; Stevens, Rowe & Badgley, 2019). These effects of history vary from mountain system and taxa, which has hampered the consolidation of a theory to explain species richness on mountain gradients (Brown, 2001). On mountaintops, (1) more geographical isolation should favor speciation and restrict processes of dispersal and colonization, and (2) species richness should decline as a consequence of higher extinction caused by generally smaller areas associated with higher elevation bands (Brown, 2001; Lomolino, 2001). This, in turn, should lead to fewer species but higher endemism on mountaintops. A recent worldwide review on plant endemism on mountains by Steinbauer et al. (2016) found a clear correlation of peak isolation with increased endemism in mountaintops. They suggested the coupling of historical climate changes with topography as a “pump” for montane endemism (Steinbauer et al., 2016). For mammals, there is growing evidence that endemism increases with elevation (Sánchez-Cordero, 2001; Heaney, 2001; Swenson et al., 2012; Chen et al., 2017), but this data is often descriptive and lacks statistical testing.

Here we surveyed non-volant small mammals across elevational gradients on two tropical mountains from the same range, Mt. Kinabalu and Mt. Tambuyukon, in northern Borneo. Mt. Kinabalu (4,095 m) is the tallest mountain in Sundaland, a tropical hotspot of biodiversity (Myers et al., 2000). A number of taxa have been surveyed across large elevational transects on this mountain: moths (Beck & Chey, 2008), ants (Brühl, Mohamed & Linsenmair, 1999; Malsch

et al., 2008), plants (Kitayama, 1992; Aiba & Kitayama, 1999; Aiba, Takyu & Kitayama, 2005; Grytnes & Beaman, 2006; Grytnes et al., 2008), oribatid mites (Hasegawa, Ito & Kitayama, 2006), snails (Liew, Schilthuizen & bin Lakim, 2010) and small mammals (Nor, 2001). These studies have recovered either a decline in diversity with elevation, which seems to fit a global pattern (Rahbek, 1995) or a mid-domain effect (MDE). To date, Nor's (2001) survey is the most comprehensive dataset of non-volant small mammals on Mt. Kinabalu. It describes a clear MDE, although the number of species he reports in the lowest elevation was unexpectedly low (only five species), compared to the maximum of approximately 35 non-volant small mammals that are known to be distributed in the lowland forests of Borneo (Nor, 2001; Payne et al., 2007). Thus, there could be a strong bias in the interpretations caused by incomplete sampling of the lowest elevation (Rahbek, 1995; Lomolino, 2001). Furthermore, Mt. Kinabalu is a hotspot of endemism for many mountain lineages (Payne et al., 2007; van der Ent, 2013; Merckx et al., 2015). The montane conditions in the upper slopes of Mt. Kinabalu are unique in Sundaland. Similar conditions are only partially found on a handful of other peaks in Borneo (ie. upper montane forest in Mt. Tambuyukon, 2,579 m, Wood & van der Ent, 2012; Mt. Trusmadi, 2,642 m, Kitayama et al., 1993; Mt. Murud, 2,423 m, Beaman & Anderson, 1997; and Mt. Mulu, 2,376 m, Collins, 1980) and on several peaks in Sumatra (Laumonier, 1997).

This study aimed to describe the effects of (1) the MDE on alpha diversity (species richness) and (2) mountain endemics on beta diversity (change in the composition of the community), for small non-volant mammals along the elevational gradient on Mt. Kinabalu and Mt. Tambuyukon. We hypothesized that alpha diversity should decrease with elevation coinciding with less available area and less complex habitat heterogeneity, while greater endemism on higher elevational bands should correlate with the particular isolation of Mt.

Kinabalu in Sundaland. For this purpose, we characterized the small mammal diversity along the elevational gradient on Mt. Kinabalu and Mt. Tambuyukon. The raw data is fully accessible and all analyses and figures are reproducible (DOI: 10.5281/zenodo.3341178).

## Materials & Methods

### *Study sites*

Mt. Kinabalu and Mt. Tambuyukon are two neighboring peaks inside Kinabalu National Park in the Malaysian state of Sabah, Borneo (Figure 1). This park covers 764 square kilometers. Mt. Kinabalu is the tallest peak in Borneo at 4,095 m and is home to thousands of endemic plant and animal species (Payne et al., 2007; van der Ent, 2013). Mt. Tambuyukon (the 3<sup>rd</sup> highest peak in Borneo, 2,579 m; Figure 1), despite being only 18 km away, is far less scientifically explored. The vegetation zones as described by Kitayama (1992) for Mt. Kinabalu have been used for simplicity as well as for consistency with previous elevational surveys (Nor, 2001): lowland (>1,200 m), lower montane (1,200-2,000 m), upper montane (2,000-2,800 m) and subalpine (2,800 -3,400 m).

The lowland dipterocarp forest dominates both mountains from the lowest elevations up to 1,200 m. Above this elevation begins the lower montane oak forest of 10-25 m trees up to around 1,800-1,900 m on both Mt. Kinabalu and Mt. Tambuyukon. On Mt. Tambuyukon at 1,440 m there is a sharp break to an ultramafic outcrop and the vegetation changes to a low productivity forest with shorter trees. The mossy or cloud forest begins at around 2,000 m on both mountains. This zone is usually immersed in clouds, moss covers most surfaces, and pitcher plants (genus *Nepenthes*), epiphytes, orchids, and climbing bamboos are abundant. At 2,350 m on Mt. Tambuyukon and 2,600 m on Mt. Kinabalu there is a fast transition to an open stunted forest dominated by *Dacrydium* and *Leptospermum* species. At these elevations, the vegetation

develops a sclerophyllous and microphyllous syndrome. At 2,800 m the subalpine vegetation appears on Mt. Kinabalu, which is absent on Mt. Tambuyukon.

### *Field survey*

Surveys were conducted in two consecutive field seasons along elevational gradients following climbing trails along Mt. Tambuyukon and Mt. Kinabalu. We targeted small non-volant mammals and further included opportunistic observations and data from trail cameras. Species identification was performed according to Payne *et al.* (2007). During the first field season, we surveyed Mt. Tambuyukon in June-August 2012. Surveys for the second field season were conducted on select locations on Mt. Tambuyukon (to fill gaps in the first season sampling) and along the full elevational gradient of Mt. Kinabalu in February-April 2013.

We set traps from ~331 - 2,509 m on Mt. Tambuyukon, and from 503 - 3,466 m on Mt. Kinabalu (Dataset S1). The taxa we expected in the small mammal trap surveys included members of the families Soricidae (shrews), Erinaceidae (gymnures), Tupaiidae (treeshrews), and rodents in the families Muridae (mice and rats) and Sciuridae (squirrels). Trapping was conducted following ethical standards according to the guidelines of the American Society of Mammalogists (Sikes et al., 2011). Animal care and use committees approved the protocols (Smithsonian Institution, National Museum of Natural History, Proposal Number 2012-04 and Estación Biológica de Doñana Proposal Number CGL2010-21524). Field research was approved by Sabah Parks (TS/PTD/5/4 Jld. 45 (33) and TS/PTD/5/4 Jld. 47 (25)), the Economic Planning Unit (100-24/1/299) and the Sabah Biodiversity Council (JKM/MBS.1000-2/2(104)).

Line transects were set at approximately every 400 - 600 m in elevation. On Mt. Tambuyukon, transects were placed along the mountaineering trail markers (placed every 1 kilometer along the trail) as follows: from Monggis substation to km 1 at 500 m, 900 m (km 7.5),



1,300 m (km 10.3), 1,600 m (km 11), 2,000 m (km 12.6) and 2,400 m (km 13.5). On Mt. Kinabalu the 500 and 900 m transects were located at Poring Hot Springs from the entrance and along the trail to the Langanan Waterfall. The next elevation transect for Mt. Kinabalu was set at ~1,500 m at the Park Headquarters, ~2,200 m along the Timpohon mountaineering trail (km 2, Kamborangoh), 2,700 m (km 4, Layang-Layang), and 3,200 m (around Waras, Pendant hut, and Panar Laban). For reporting results and data analysis, we hereafter refer to these locations as “trapping locations”. Each trapping location gathered the trap data from transects which could span a distance of over 100 m up- or downhill (raw data in Dataset S1).

We set traps at approximately 5 - 10 m intervals for a total of around 40 traps per transect. Trapping locations are shown in Figure 1. Collapsible Tomahawk live traps (40 cm long), collapsible Sherman traps (two sizes used: 30 cm and 37 cm long), and local mesh-wire box traps were used. We considered traps as ‘close to the ground’ if set below approximately 3 m off the ground. Most of these traps were directly set on the ground, while a small proportion was set on branches or vines at the reach of the hand. We considered the 3-m threshold as reasonable to describe the terrestrial small mammal community given the arboreal habits of many small terrestrial Bornean mammals (Wells et al., 2004) and the high-complexity of the vegetation in most transects. Any trap above that was considered ‘arboreal’. A bait mixture (of varying composition) consisting of bananas, coconuts, sweet potatoes, palm fruit and oil, vanilla extract, and dried fish was placed in each trap. A small number of pitfall traps were distributed from 500 - 2,000 m on Mt. Tambuyukon (number and distribution were limited by the substrate) (Table 1).

Each trapping location had a total of 2 - 4 transects. The highest elevation had a lower number of trap nights due to the smaller area available for placement of traps. Coordinates for trapping locations were recorded using Garmin eTrex® series and Garmin GPSmap 60CSx. The

minimum number of trap nights was based on the saturation rates obtained from Nor (2001) at approximately 300 trap nights. For every sampling location (elevation within mountain) the cumulative trap-nights and species richness were calculated by adding the cumulative number of species caught and trap-nights for each trap for each successive day from the date the trap was set.

We set up 4 camera traps (Reconyx RapidFire RC55™ cameras, and ScoutGuard HCO™ cameras) along the mountaineering trail on Mt. Tambuyukon. Camera 1 was placed at 500 m, at the first-kilometer marker for the hiking trail. Cameras 2 and 3 were placed along the Kepuakan River near km 8 and at approximately 900 m. Camera 4 was placed at approximately 1,300 m near km 10.5. No cameras were deployed along the Mt. Kinabalu trail due to a large number of day hikers and mountain climbers.

Additionally, while on Mt. Tambuyukon we opportunistically recorded mammal observations while walking to, from and along our trap lines, while setting cameras, or while in our campsite.

# *Alpha diversity*

*Diversity index.* We computed diversity indices for each trapping location. The Community Ecology Package ‘vegan’ 2.5-5 (Dixon, 2003) in R 3.5.1 (R Development Core Team, 2018) was used to calculate the Shannon diversity ( $H'$ ) and Simpson’s diversity. Pielou’s evenness ( $J'$ ) was calculated as  $J' = H' / H_{\max}$ , and species richness ( $S$ ) as the number of species. We used the LOWESS smoother (*stats::lowess* function, in R) to visualize the change of these indexes with elevation. The pattern of species richness across elevation in each mountain was contrasted against the mid-domain effect using ‘rangemodelR’ 1.0.4 (Marathe, 2019) in R, as in Wang & Fang, 2012. We used the function *range.shuffle* which returns the pattern of species

diversity under the mid-domain effect. The following arguments were used for computations: 50  
m as the interval to discretize species ranges and midpoints into elevational bands, soft  
boundaries, and 5,000 repetitions.

*Predictors of species richness.* The relation between species richness and two predictors,  
(1) elevation and (2) expectations under the MDE, were assessed in separate linear models with  
mixed-effects (Laird & Ware, 1982) using *lme* function from ‘nlme’ package 3.1-137 (Pinheiro  
et al., 2018) in R. Model building and the evaluation of their fit was carried following the  
recommendations in Harrison et al. (2018). In both models, we corrected for spatial  
autocorrelation by including a term to account for the correlation structure built with  
*nlme::corSpatial* function in R. This object contained the centroids of coordinates from all  
animals trapped at a given location. Mountain was included as a random intercept. The fit of  
these models was compared using likelihood ratio tests (King, 2010) against a null model that  
excluded the fixed effect (Nickerson, 2000; Pinheiro & Bates, 2006; Harrison et al., 2018).

*Endemism.* We quantified endemism at each trapping location as the proportion of  
endemic species. This measure is robust to sampling bias and it is not overridden by local  
diversity (Steinbauer et al., 2016). We defined endemic as only found in Borneo. Shrews were  
excluded as we were not able to identify them to the species level. To visualize the change of  
proportion of endemics with elevation we created a confidence envelope by resampling 1,000  
times the species present at each trapping location. This same approach was used to visualize the  
proportion of captures that belonged to endemic species across elevation. We evaluated the  
contribution of elevation (predictor) to explaining the proportion of endemism (dependent  
variable 1) and the proportion of captures (dependent variable 2) that corresponded to endemic  
species. We used generalized models with mixed effects using a binomial family with logit link.

Mountain was added as a random intercept. Model fitting was done using *glmer* function from ‘lme4’ package 1.1-21 (Bates et al., 2015) in R. The fit of these models was compared using likelihood ratio tests against a null model that excluded the fixed effect (Nickerson, 2000; Pinheiro & Bates, 2006).

# *Beta diversity*

We calculated beta diversity for each mountain using a Sorensen-based dissimilarity index ( $\beta_{\text{SOR}}$ ) with its turnover ( $\beta_{\text{SIM}}$ ) and nestedness ( $\beta_{\text{NES}}$ ) components (Baselga, 2010). For calculations we used function *beta.multi*, from package ‘betapart’ 1.3 (Baselga & Orme, 2012) in R. These calculations were also done pairwise between all trapping locations within the same mountain using the function *betapart::beta.pair*. These distances were used for a cluster analysis using neighbor-joining with (Saitou & Nei, 1987) using *ape::nj* function in R to evaluate the community similarity between mountains and elevations.

We estimated the contribution of endemics to  $\beta_{\text{SIM}}$  and  $\beta_{\text{NES}}$  components of beta diversity. For each mountain, we removed endemic species from the dataset and recalculated  $\beta_{\text{SIM}}$  and  $\beta_{\text{NES}}$  (hereafter referred to as  $\beta_{\text{SIM-end}}$  and  $\beta_{\text{NES-end}}$ ). We created subsets of the data to which these measures could be directly compared by randomly sub-sampling 5,000 times without replacement, several species from the complete dataset equal to the non-endemics present in that mountain. P-values were determined from the distributions of the permuted values.

## 242 Results

### 243 *Field survey*

244 The overall trapping success increased towards high elevation on both mountains. On Mt.  
 245 Tambuyukon, we trapped a total of 295 different individuals (not including recaptured animals)  
 246 from 21 different species (Dataset S1, Table S1) over 5,957 trap nights, for a total of 5.0% trap  
 247 success (not including arboreal or pitfall trapping; Table 1). Trap success at each elevation  
 248 ranged from 2.1% at 1,600 m to 9.6% at 2,400 m. One non-target capture of a carnivore, a  
 249 Kinabalu ferret-badger (*Melogale everetti*), was recorded. The trap success calculations were  
 250 done excluding pitfall traps and arboreal traps (due to inconsistent placement of traps). The use  
 251 of pitfall and arboreal traps was limited by the time necessary to set and check arboreal traps,  
 252 difficulty finding appropriate places to set pitfall traps and the high number of tourists on Mt.  
 253 Kinabalu, which limited the sites we could set traps to those out of sight of the trails. The  
 254 accumulation of species across trap nights varied across elevations, and appeared near saturation  
 255 in all trapping locations (Figure S1). A species of shrew, *Suncus sp.*, was collected in a pitfall  
 256 trap, and a gray tree rat (*Lenothrix canus*) in an arboreal trap, bringing the total number of  
 257 species to 23.

258 On Mt. Kinabalu, we trapped a total of 20 species from 209 different individuals (Dataset  
 259 S1; Table S1) over 2,022 trap nights, for an average trap success of 10.3% (Table 1). The trap  
 260 success across elevations was much higher on Mt. Kinabalu, ranging from 5.6% (at 900 m), to  
 261 15.4% (at 2,700 m) (Table 1). This overall higher capture rate resulted in species saturation with  
 262 a lower number of trap nights on Mt. Kinabalu than on Mt. Tambuyukon (Figure S1).

# Species distribution

The mountain treeshrew, *Tupaia montana*, was the most frequently caught species (35.7 % of all catches) and it had a wide elevational distribution from 836 - 3,382 m. The Bornean mountain ground squirrel (*Sundasciurus everetti*, formerly *Dremomys everetti*; Hawkins et al. 2016), the long-tailed giant rat (*Leopoldamys sabanus*), and Whitehead's spiny rat (*Maxomys whiteheadi*) also had large elevational distributions on both mountains (Figure 2). The lowland (< 1,000 m) terrestrial small mammal community was the most diverse with 19 species trapped. We captured 16 species in the community associated with montane forest between 1,000 - 2,400 m and only 7 species at 2,400 m and above (Figure 2).

We captured a single shrew (*Suncus sp.*) after a trapping effort of 176 pitfall trap nights. A less intensive arboreal trapping effort of 76 trap nights yielded seven individuals from six small mammal species: *Tupaia montana* (n = 2), *Lenothrix canus* (n = 1), *Callosciurus prevostii* (n = 1), *Sundasciurus jentinki* (n = 1), *Sundamys muelleri* (n = 1) and *Tupaia minor* (n = 1). Despite the smaller effort of arboreal trap nights, we still captured two species that were not trapped elsewhere (*Lenothrix canus* and *Sundasciurus jentinki*). We used a threshold of three meters off the ground to consider a trap arboreal. However, we also caught arboreal species in ground traps or traps close to the ground (< 3m), including *Chiropodomys pusillus*, *Callosciurus prevostii* and *Tupaia minor*.

We set 4 trail cameras on Mt. Tambuyukon to document larger mammals not targeted by our traps. They documented an additional 8 species of mammals (Table 2; Figure S2). The number of species captured by the cameras varied from one to five, with the camera at 500 m exhibiting the most diversity, both in number of species and number of independent visits (Table 2). They documented four species that were not documented in any other way.

On Mt. Tambuyukon, several species were detected only through direct observation (Table 3). Of these sightings many were documented only a single time, including the orangutan (*Pongo pygmaeus*), the Bornean giant tufted ground squirrel (*Reithrosciurus macrotis*), Whitehead's squirrel (*Exilisciurus whiteheadii*), and the bearded pig (*Sus barbatus*). The Bornean gibbon (*Hylobates muelleri*) was heard singing on an almost daily basis, but only directly observed a single time. The sambar deer (*Rusa unicolor*) was heard vocalizing once at 1,400 m. Only one observation was made of a carnivore, the Malay civet (*Viverra zibetha*), which was observed during a late night-walk. The visual observations increased the diversity of species documented, especially for primates and tree squirrels.

#### *Alpha diversity*

Both mountains showed a similar pattern for alpha diversity indices across elevations (Figure 3; Table S2). Species richness and Shannon diversity were maximum in low elevations and decreased gradually towards high elevations. However, evenness was lowest at middle elevations (U-shaped) (Figure 3). The high dominance of some species at middle elevations (e.g. mountain treeshrew) leads the Shannon diversity to sink at around 1,500 m in both mountains. However, Shannon diversity increases again towards the highest elevations due to the more even occurrence of the species in the small mammal communities at these elevations, despite the lower species richness (Figure 3). The species richness we report in low elevations for both mountains are above the upper 97.5% quantile of the expected richness expected under the hypothesis of the mid-domain effect (Figure 3A).

We detected a significant negative relationship between species richness and elevation ( $\chi^2_1 = 8.81$ ,  $p = 0.003$ ). However, species richness was not correlated with expected species richness under the MDE ( $\chi^2_1 = 0.54$ ,  $p = 0.46$ ) (Figure 3).

Elevation was positively correlated with higher proportion of endemic species ( $\chi^2 = 7.96$ ,  $df = 1$ ,  $p = 0.005$ ) and a greater proportion endemic species captured ( $\chi^2 = 10.7$ ,  $df = 1$ ,  $p = 0.001$ ) (Figure 4).

# *Beta diversity*

Variation in the species composition assemblages, or beta diversity, was very similar for both mountains ( $\beta_{SOR} = 0.77$  on Mt. Kinabalu and  $\beta_{SOR} = 0.74$  on Mt. Tambuyukon). Most of this beta diversity derived from the turnover component ( $\beta_{SIM} = 0.73$  for Mt. Kinabalu and 0.65 for Mt. Tambuyukon). The nestedness component was very low on both mountains ( $\beta_{NES} = 0.04$  for Mt. Kinabalu and 0.09 for Mt. Tambuyukon) (Figure 5). This indicates that the assemblages at different elevations are not the product of species loss from the richest assemblages. Instead, they are singular assemblages with different species compositions. We found large dissimilarities in the turnover component between pairwise locations within each mountain ( $\beta_{sim}$ ) associated with  $< \sim 1900$  m vs  $> \sim 1900$  m locations, compared to lower values within lowland or highland locations (Tables S3). When removing endemic species from the dataset,  $\beta_{NES-end}$  increased and  $\beta_{SIM-end}$  decreased on both mountains (Figure 5). The permutations indicated that this decrease in the turnover component ( $\beta_{SIM-end}$ ) was significant in Tambuyukon ( $\beta_{SIM-end} = 0.36$ ,  $p = 0.002$ ), but not in Kinabalu ( $\beta_{SIM-end} = 0.68$ ,  $p = 0.36$ ), and that the increase in nestedness ( $\beta_{NES-end}$ ) was significant for Mt. Tambuyukon ( $\beta_{NES-end} = 0.38$ ,  $p = 0.001$ ), but only marginally significant for Mt. Kinabalu ( $\beta_{NES-end} = 0.14$ ,  $p = 0.05$ ) (Figure 5).

The clustering analysis grouped trapping locations in 2 main groups, above and below  $\sim 1,400$  m (Figure S3). Within the high elevation group, all locations above  $\sim 1,900$  m clustered together.



# Discussion

We report an extensive survey of non-volant small mammals within Kinabalu National Park, Borneo. A sampling effort of 8,231 trap-nights in Mt. Kinabalu (n = 2,044) and its neighbor peak Mt. Tambuyukon (n = 6,187) yielded a total of 512 individual records (Mt. Kinabalu, n = 213; Mt. Tambuyukon, n = 299) from 27 species (Mt. Kinabalu, n = 20; Mt. Tambuyukon, n = 23) (Table 1; Figure 2). Records from camera traps and direct observations increased the total number of species recorded for Mt. Tambuyukon by 18 to a total of 44 (Table 3).

Mt. Kinabalu is a biodiversity hotspot for many taxa (van der Ent 2013). Its mammal fauna has been studied for over a century (eg. Oldfield, 1889; Whitehead, 1893; Emmons, 2000; Nor, 2001) and it is known to host 61 species of non-volant small mammals (Nor, 2001; Payne et al., 2007). We further explored for the first time the non-volant small mammal diversity along the complete elevational gradient in Mt. Tambuyukon (2,579 m), the third-highest peak in Borneo and only 18 km away from Mt. Kinabalu. Some important sightings on Mt. Tambuyukon included the orangutan (*Pongo pygmaeus*), which has an estimated population of only 50 individuals within Kinabalu Park boundaries (Ancrenaz et al., 2005). The Kinabalu ferret-badger (*Melogale everetti*) was a significant finding since it is the first official record of this species on Mt. Tambuyukon (Payne et al., 2007; Wilting et al., 2016). We trapped this species at 2,051 m on Mt. Tambuyukon and 3,336 m on Mt. Kinabalu. We identified a population of the summit rat (*Rattus baluensis*) on Mt. Tambuyukon, previously only known from Mt. Kinabalu. This species was common at high elevations and has its lower distribution limit at around 2,000 m. A population genetic analysis of the summit rats from Mt. Kinabalu and Mt. Tambuyukon demonstrated that they are currently genetically isolated (Camacho-Sanchez et al. 2018). We

also make the first records of the mountain species *Maxomys alticola*, *Melogale everetti*, *Hylomys suillus* and *Niviventer rapit* on Mt. Tambuyukon.

*Alpha diversity*

Species richness peaked at low elevations on both mountains coinciding with the lowland dipterocarp forest. Then, it decreased gradually towards the highest elevations where it was lowest (Figure 3). This pattern deviates from the expectations of the MDE (Figure 3), and the MDE reported by Nor (2001) in a previous small mammal survey on Mt. Kinabalu with a very similar survey scheme to ours. Colwell & Lees (2000) suggested that a MDE should constitute the null hypothesis over which deviations should be interpreted. However, this point of view is not universal (Rahbek, 1995; McCain, 2007, 2009). Incomplete sampling of the low elevations happens regularly and can artificially create a MDE (Rahbek, 1995; Lomolino, 2001), which may explain the difference between Nor (2001) and our results for the same mountain. Our surveys detected 12 more species as compared to Nor (2001), including a climbing mouse, *Chiropodomys pusillus*, a tree rat *Lenothrix canus*, Prevost's squirrel, *Callosciurus prevostii*, Jentink's squirrel, *Sundasciurus jentinki*, the rats *Maxomys rajah*, *M. alticola* and *Sundamys muelleri*, the non-native species *Rattus exulans* and *R. tiomanicus*, two species of treeshrews *Tupaia longipes* and *T. minor*, and two species of shrews, one trapped in a small Sherman trap, *Crocidura* sp., and one in a pitfall trap, *Suncus* sp.. The effects of an incomplete sampling should be more acute in lowland elevations where there is more habitat heterogeneity (Rosenzweig, 1992, 1995) and species might tend to occupy smaller ranges (Rosenzweig, 1995; Brown, 2014). However, we report the highest alpha diversity in the lowest elevations, which supports that our sampling scheme was not affected by this low-elevation sampling bias. Another observation supporting the comprehensiveness of our sampling is that we documented species in all sites

between the lowest and highest occurrences, except for *M. whiteheadi* on Mt. Kinabalu, and *M. ochraceiventer* and *N. rapit* on Mt. Tambuyukon.

Conversely, the gradual decrease of species richness we recorded was explained by elevation alone. Even in the presence of a MDE, a gradual decrease of species richness with elevation seems to be a general pattern in mountain gradients (Rahbek, 1995). There are multiple factors that are correlated with elevation which have been proposed to explain diversity across mountains gradients, but disentangling their effects is difficult given the multicollinearity (Heaney, 2001; Fu et al., 2006; Kluge, Kessler & Dunn, 2006). The strong correlation of alpha diversity with elevation enables further discussion. Diversity has been proposed to peak with precipitation (Heaney, 2001), but on Mt. Kinabalu the peak of precipitation, at around 2,000 m (Kitayama, 1992), did not match the diversity peak. Productivity (Aiba, Takyu & Kitayama, 2005) and temperature (Kitayama, 1992) are negatively correlated with elevation in Kinabalu, and could potentially explain the diversity pattern. Nevertheless, it has been suggested that examining resource availability for this taxonomic group is more appropriate than simply looking at productivity (Brown, 2001; Heaney, 2001). However, this is a difficult variable to measure that was not incorporated in our original survey experimental design. A more plausible explanation for the change in diversity is area and habitat complexity. Available area (Camacho-Sanchez et al., 2018) and complexity of the forest (Kitayama, 1992) decrease with elevation on Mt. Kinabalu. The peak in diversity we find in low elevations is consistent with the spatial hypothesis which states that (1) at the regional level, larger areas (such as the lower elevations in mountains) have lower rates of extinction over speciation (Rosenzweig, 1992) and (2) that larger areas have more types of different habitats, so greater species diversity should be observed in larger areas (Rosenzweig, 1995). Perhaps, the relationship between area and diversity on

elevational gradients along large mountains falls somewhere between these processes (McCain, 2007).

# *Beta diversity*

The composition of the small mammal assemblage changed across elevation in a similar way between mountains. Indeed, all trapping locations in high elevations (above ~1,900 m) were very similar in composition (Figure S3). Mid-mountain locations showed intermediate compositions whereas the 500 m and 900 m locations on both mountains also clustered together. This montane fauna transition was already identified by Nor (2001) to be at around 1,800 m, and it matches approximately the vegetation limit between the lower and upper montane forests on Mt. Kinabalu (Kitayama 1992). The influence of this shift is reflected in the overall high turnover component of beta diversity ( $\beta_{SIM}$ ) for both mountains (Figure 5). The pairwise turnover components ( $\beta_{sim}$ ) were highest between lowland - highland locations (Tables S3). This indicates the lowland and highland communities are composed of different species, rather than the community with the lowest richness (highland) being a subset of the species present on the richest one (lowland). This pattern has already been described for other small mammals across several mountain systems (Mena & Vázquez-Domínguez, 2005). Unfortunately, there is no consistent data collection from other tall mountains in Sundaland to discuss a common mountain biogeography pattern in this region. Historical expeditions to Sumatra also point to a similar shift in vegetation structure and different mammal assemblages at high elevations above 2,000 m (Robinson & Kloss, 1918, 1919; Miller, 1942).

# *Pattern of endemism*

Beyond the net alpha and beta diversity we describe, we place special attention on the nature of the species that could be driving these diversity patterns. Our models predicted that the

proportion of Bornean endemism as well as the proportion of endemic species captured increased with elevation (Figure 4). At the same time, we found that the high-elevation endemics were responsible for the high turnover component in beta diversity in both mountains (Figure 5). The species that had the greatest contribution to this high-elevation endemism were *Rattus baluensis*, *Tupaia montana*, *Sundasciurus everetti* and *Maxomys alticola* (Table S1). These species are restricted to mountain areas in northern or central-northern Borneo, and were present in high abundance and evenness from 2,000 m (Table S1). Nor (2001) reported high trapping success on Mt. Kinabalu, associated with higher abundance at these elevations, but he did not record *Maxomys alticola*. Heaney (2001) also recorded the highest abundances in the top elevations in the Philippines, and a peak of endemics at higher elevations. Three additional trapped species are mountain endemics (*Melogale everetti*, *Sundasciurus jentinki* and *Niviventer rapit*) but contributed less to our analysis because of their lower densities or detectability. This pattern of high-elevation endemics could be even more pronounced as the distribution and taxonomy of the highland mammals also found on other islands in Sundaland are further updated and revised (i.e. *Hylomys suillus* and *Sundamys infraluteus*; Camacho Sánchez, 2017). A majority of the lowland species are widespread, and also distributed across other Sundaland landmasses such as Sumatra and the Malay Peninsula (Corbet & Hill, 1992).

The pattern of mountain endemics on Mt. Kinabalu has previously been described for other taxa (Merckx et al., 2015). For mammals, this has also been observed by Heaney (2001), Sanchez-Cordero (2001), Swenson et al. (2012) and Chen et al. (2017) on other mountains. Mt Kinabalu is unique in Sundaland due to its high elevation (4,095 m) and the scarcity of nearby peaks above 2,000 m. Higher isolation on mountain peaks boosts mountain endemism worldwide (Steinbauer et al., 2016), which could explain the greater endemism in the higher elevations in

the Kinabalu range. For instance, a pattern of mountain endemism linked to divergence in allopatry induced by isolation from the combination of topography with past climate changes has been described in Bornean birds (Sheldon, Lim & Moyle, 2015; Moyle et al., 2017; Manthey et al., 2017). A similar pattern of high mountain endemism driven by intra-island speciation has been reported for shrews in Sumatra and Java (Esselstyn et al., 2013; Demos et al., 2016). Presumably, the high degree of isolation of the habitats on the higher slopes of Mt. Kinabalu help to maintain a highly endemic community. This could be due to reduced dispersal and colonization to/from nearby similar areas as proposed by Steinbauer et al. (2016).

## Conclusions

We found a decline in small mammal diversity from low to high elevations on both Mt. Kinabalu and Mt. Tambuyukon. This pattern differs from the MDE previously described for Mt. Kinabalu and other mountains worldwide. The decrease in diversity with elevation is concordant with the spatial hypothesis predicting higher diversity in lowlands driven by historically larger areas with less extinction and more habitat heterogeneity. However, we cannot exclude other climatic or ecological hypothesis. Endemic species were in higher proportion and more abundant in higher elevations and they drove the turnover component of beta diversity. The high number of mountain endemics point to historical factors as important drivers of the biogeography in this region.

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# **Supplementary Information**

Dataset S1: Data on animals sampled and trapping effort.  
Table S1: Summary table of species per elevation.  
Figure S1: Trap-night accumulation curves  
Figure S2: Photographs from camera traps.  
Table S2: Diversity indices.  
Figure S3: Cluster analysis.  
Tables S3: Pairwise beta diversity.

**Table 1**(on next page)

Trap success across all elevations.

Trap success across all elevations. The number of animals caught is in column N, followed by number of trap nights, and the overall trap success per altitude.

Table 1: Trap success across all elevations. The number of animals caught is in column N, followed by number of trap nights, and the overall trap success per altitude.

	Elev. (m)	Including arboreal and pitfall traps			Excluding arboreal and pitfall traps		
		N	Trap nights	Trap success	N	Trap nights	Trap success
Kinabalu	500	33	300	11.0%	30	285	10.5%
	900	20	360	5.6%	20	360	5.6%
	1,500	36	360	10.0%	36	360	10.0%
	2,200	35	434	8.1%	34	427	8.0%
	2,700	60	390	15.4%	60	390	15.4%
	3,200	29	200	14.5%	29	200	14.5%
	Totals	213	2,044	10.4%	209	2,022	10.3%
Tambuyukon	500	78	1713	4.6%	75	1588	4.7%
	900	24	992	2.4%	24	956	2.5%
	1,300	53	712	7.4%	52	702	7.4%
	1,600	22	1036	2.1%	22	1,025	2.1%
	2,000	55	1036	5.3%	55	988	5.6%
	2,400	67	698	9.6%	67	698	9.6%
	Totals	299	6,187	4.8%	295	5,957	5.0%

## **Table 2**(on next page)

Results of camera trap survey.

Results of camera trap surveys on Mt. Tambuyukon, with relative abundance calculated for 100 trap nights.

1 Table 2: Results of camera trap surveys on Mt. Tambuyukon, with relative abundance calculated  
2 for 100 trap nights.

Camera	Elevation (m)	Common Name	Species	No. of series	Camera nights	Relative abundance
1	500	Pig-tailed Macaque	<i>Macaca nemestrina</i>	1	42	2.38
		Common Porcupine	<i>Hystrix brachyura</i>	2		4.76
		Mouse Deer	<i>Tragulus</i> sp.	2		4.76
		Muntjac	<i>Muntiacus</i> sp.	1		2.38
		Sambar Deer	<i>Rusa unicolor</i>	1		2.38
2	900	Malay Civet	<i>Viverra zibetha</i>	2	42	9.52
		Banded Linsang	<i>Prionodon linsang</i>	1		2.38
3	900	Malay Civet	<i>Viverra zibetha</i>	2		----
4	1,300	Malay Civet	<i>Viverra zibetha</i>	1	29	3.45
		Masked Palm Civet	<i>Paguma larvata</i>	1		3.45

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**Table 3**(on next page)

All species recorded on Mt. Tambuyukon.

All species recorded on Mt. Tambuyukon.

1 Table 3: All species recorded on Mt. Tambuyukon.

Number	Family	Common name	Scientific name	Method(s) of detection
1	Cercopithecidae	Pig-tailed Macaque	<i>Macaca nemestrina</i>	camera trap/ observation
2	Cercopithecidae	Long tailed Macaque	<i>Macaca fascicularis</i>	observation
3	Cercopithecidae	Maroon Langur	<i>Presbytis rubicunda</i>	observation
4	Cervidae	Muntjac	<i>Muntiacus</i> sp.	camera trap
5	Cervidae	Sambar Deer	<i>Cervus unicolor</i>	camera trap/ observation
6	Erinaceidae	Lesser Gymnure	<i>Hylomys suillus</i>	live trap
7	Hylobatidae	Bornean Gibbon	<i>Hylobates muelleri</i>	observation
8	Hystriidae	Common Porcupine	<i>Hystrix brachyura</i>	camera trap
9	Muridae	Common Pencil-tailed Tree Mouse	<i>Chiropodomys pusillus</i>	live trap
10	Muridae	Grey tree rat/ Sundaic Lenothrix	<i>Lenothrix canus</i>	live trap
11	Muridae	Long-tailed giant rat	<i>Leopoldomys sabanus</i>	live trap
12	Muridae	Bornean Mountain Maxomys	<i>Maxomys alticola</i>	live trap
13	Muridae	Chestnut-bellied spiny rat	<i>Maxomys ochraceiventer</i>	live trap
14	Muridae	Brown Spiny Rat	<i>Maxomys rajah</i>	live trap
15	Muridae	Red Spiny Rat	<i>Maxomys surifer</i>	live trap
16	Muridae	Whitehead's Rat	<i>Maxomys whiteheadi</i>	live trap
18	Muridae	Dark-tailed tree rat	<i>Niviventer cremioventer</i>	live trap
19	Muridae	Mountain long tailed rat	<i>Niviventer rapit</i>	live trap
20	Muridae	Summit Rat	<i>Rattus baluensis</i>	live trap
21	Muridae	Polynesian/Pacific rat	<i>Rattus exulans</i>	live trap
22	Muridae	Giant Mountain Rat	<i>Sundamys infraluteus</i>	live trap
23	Muridae	Muller's Rat/ Sundamys	<i>Sundamys muelleri</i>	live trap
24	Mustelidae	Kinabalu ferret-badger	<i>Melogale everetti</i>	live trap
25	Pongidae	Bornean Orangutan	<i>Pongo pygmaeus</i>	observation
26	Sciuridae	Bornean Mountain Ground Squirrel	<i>Sundasciurus everetti</i>	live trap
27	Sciuridae	Low's squirrel	<i>Sundasciurus lowii</i>	live trap
28	Sciuridae	Plantain Squirrel	<i>Callosciurus notatus</i>	observation
29	Sciuridae	Kinabalu Squirrel	<i>Callosciurus baluensis</i>	observation
30	Sciuridae	Giant Squirrel	<i>Ratufa affinis</i>	observation
31	Sciuridae	Jentink's Squirrel	<i>Sundasciurus jentinki</i>	live trap

32	Sciuridae	Whitehead's Squirrel	<i>Exilisciurus whiteheadi</i>	observation
33	Sciuridae	Giant Bornean Tufted Ground Squirrel	<i>Reithrosciurus macrotis</i>	observation
34	Soricidae	Shrew	<i>Crocidura</i> sp.	live trap
35	Soricidae	Shrew	<i>Suncus</i> sp.	live trap
36	Suidae	Bearded Pig	<i>Sus barbatus</i>	observation
37	Tragulidae	Mouse Deer	<i>Tragulus</i> sp.	camera trap
38	Tupaiaidae	Common treeshrew	<i>Tupaia longipes</i>	live trap
39	Tupaiaidae	Lesser treeshrew	<i>Tupaia minor</i>	live trap
40	Tupaiaidae	Mountain treeshrew	<i>Tupaia montana</i>	live trap
41	Tupaiaidae	Large treeshrew	<i>Tupaia tana</i>	live trap
42	Viverridae	Malay Civet	<i>Viverra zibetha</i>	camera trap
43	Viverridae	Banded Linsang	<i>Prionodon linsang</i>	camera trap
44	Viverridae	Masked Palm Civet	<i>Paguma larvata</i>	camera trap

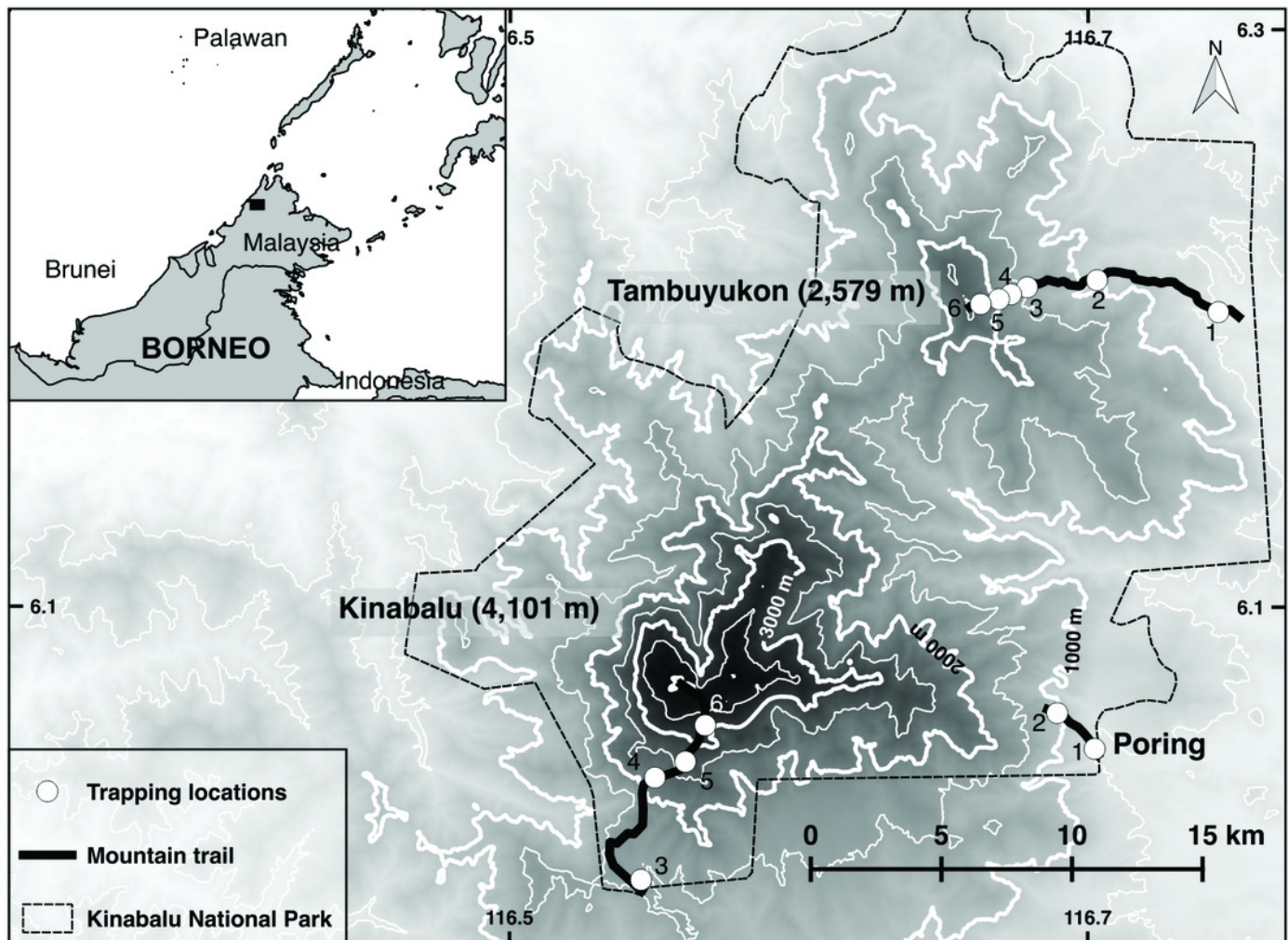
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# Figure 1

Trapping locations

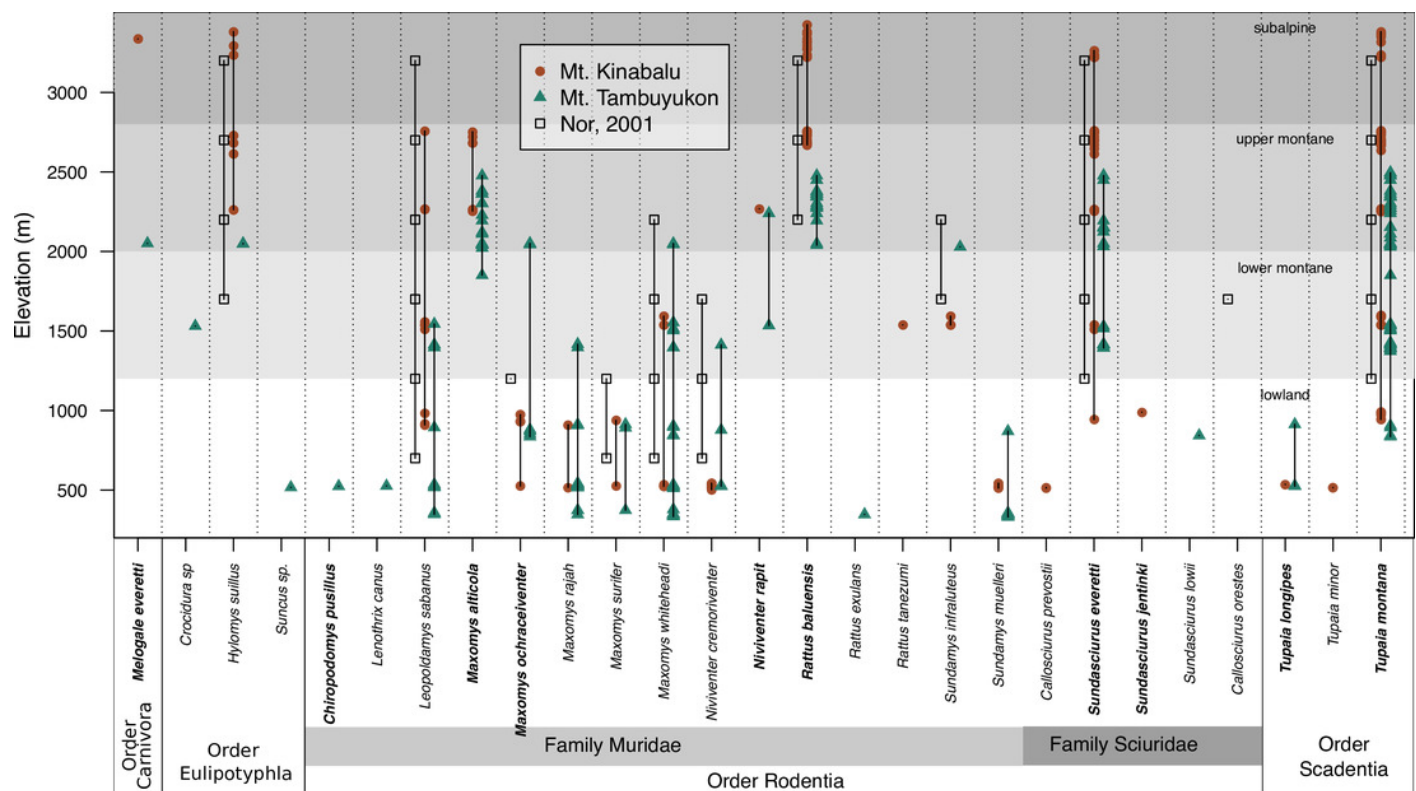
Map of Kinabalu Park, Sabah, Malaysia with trails followed and trapping locations



# Figure 2

Species distribution across elevations.

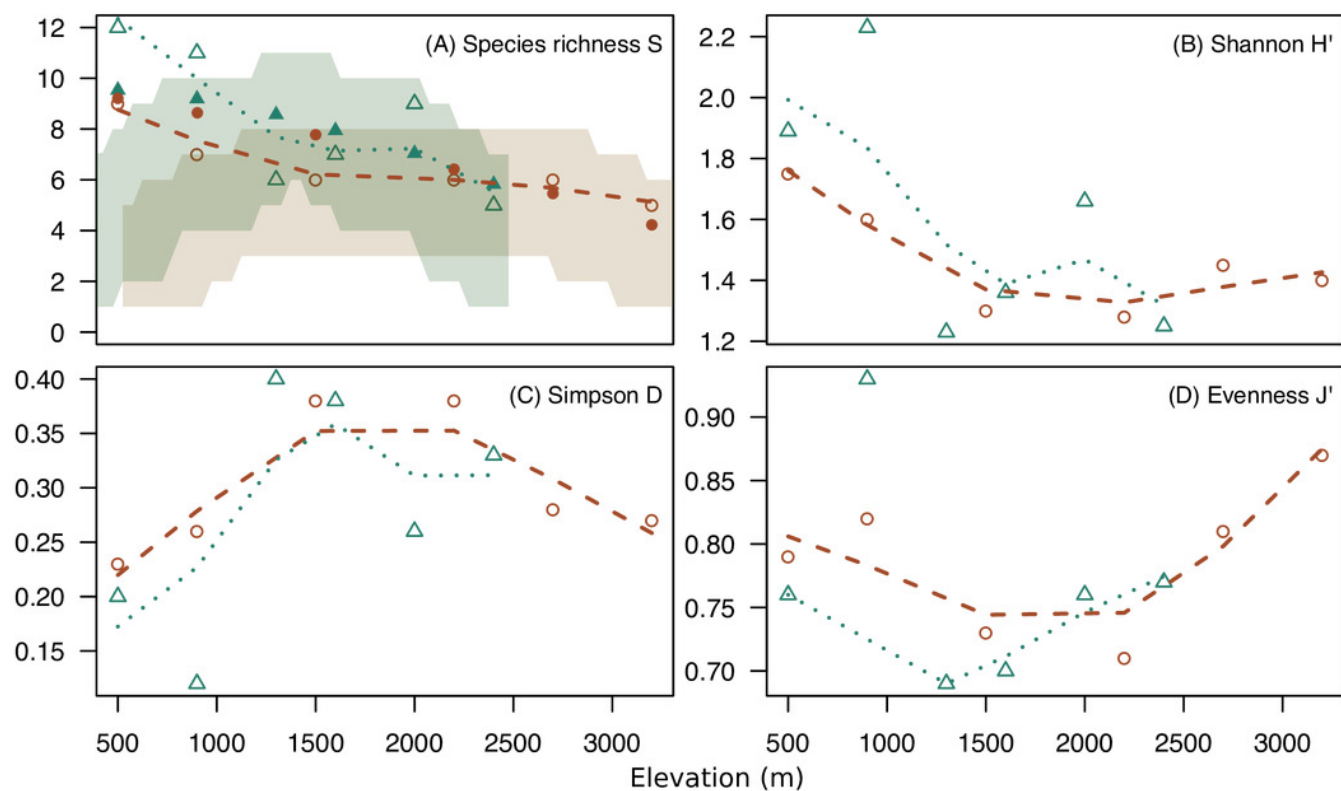
Species distribution across elevations. Our two field surveys are represented by circles (Mt. Kinabalu), and triangles (Mt. Tambuyukon), together with a previous small mammal survey on Mt. Kinabalu (open squares; Nor, 2001). Endemic species are bolded. We have represented the vegetation levels as described in Kitayama (1992) for Mt. Kinabalu in greyscale in the background.



# Figure 3

Diversity indices across elevations.

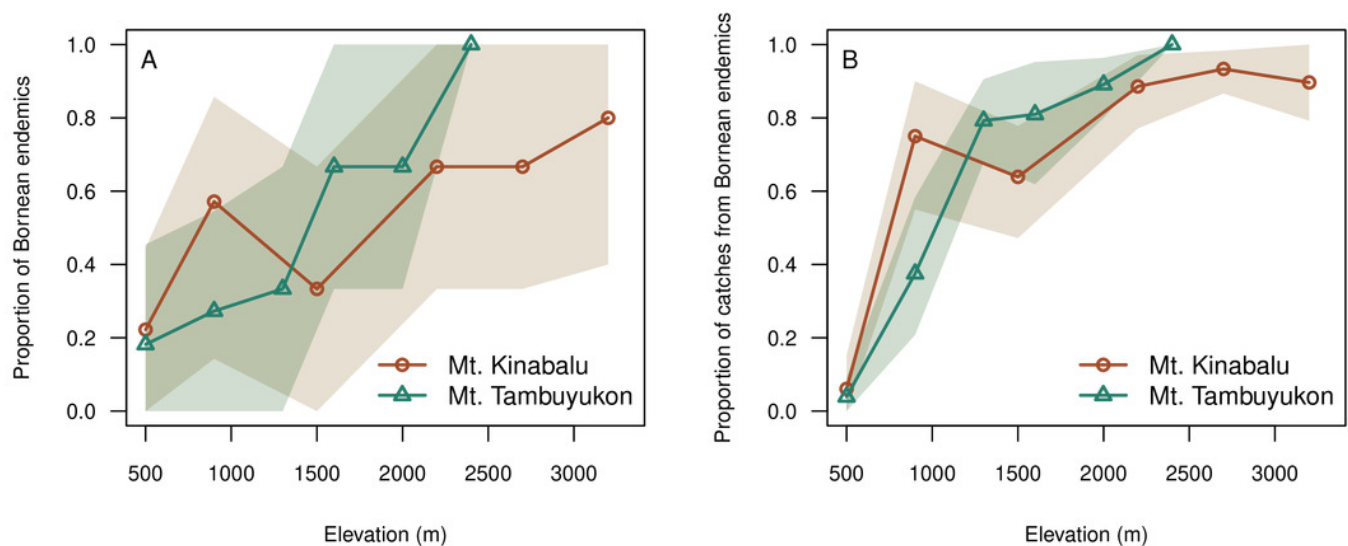
Diversity indices across elevations for Mt. Kinabalu (circles) and Mt. Tambuyukon (triangles) with loess regressions (Mt. Kinabalu, dashed line; Mt. Tambuyukon, pointed line). Shaded areas represent the 2.5% and 97.5% percentiles of the species richness for expectations under the MDE (Mt. Kinabalu, brown; Mt. Tambuyukon, green). Closed symbols in panel A are the fitted values of species richness corrected for autocorrelation.



# Figure 4

## Endemism with elevation

Proportion of Bornean endemics (A) and proportion of catches belonging to Bornean endemics (B) across elevation. A confidence envelope for the observed values is represented as a shaded area from 1000 bootstrap replicates.





# Figure 5

## Beta diversity with and without endemics

Sorensen dissimilarity ( $B_{SOR}$ ) decomposed in its nestedness ( $B_{SNE}$ ) and turnover ( $B_{SIM}$ ) components, for Mt. Kinabalu and Mt. Tambuyukon (left/right depiction in set of data points, respectively). Solid horizontal lines represent the observed values, while the dotted horizontal lines are the estimated beta diversity measures after removing endemic species from the matrix ( $B_{SIM-end}$ ,  $B_{SNE-end}$  and  $B_{SOR-end}$ ). The random expectations for  $B_{SIM-end}$ ,  $B_{SNE-end}$  and  $B_{SOR-end}$  are represented from the dotted cloud with 5,000 permuted values (see main text), together with their corresponding 2.5% and 97.5% percentiles (vertical grey bar).

