



RESEARCH ARTICLE - ANTS

Ant Species Distribution Models Reveal Mixed Elevational Diversity Patterns in Complete Elevation Gradients: Past and Present Climatic Effects?

SONIA I. PAPAQUI-BELLO, ANDRÉS LIRA-NORIEGA, ROGER GUEVARA

INECOL – Instituto de Ecología, Xalapa, Veracruz, Mexico

Article History

Edited by

Kleber Del-Claro, UFU, Brazil

Received 27 February 2024

Initial acceptance 29 February 2024

Final acceptance 25 April 2024

Publication date 25 June 2024

Keywords

Alpha diversity, climatic stability, elevational range shifts, environmental gradients, beta diversity, ecological niche models, Formicidae.

Corresponding author

Roger Guevara

INECOL – Instituto de Ecología

Xalapa, Veracruz, Mexico.

E-Mail: roger.guevara@inecol.mx

Abstract

The juxtaposition of climatic zones in tropical mountains is considered one of the leading causes of the high diversity of tropical mountains; high peaks block oceanic winds, causing mid-elevation peaks of precipitation and diversity. Also, historically wind-protected valleys may harbor high species richness because of climate stability. Despite this knowledge, the elevational patterns of many taxa and their drivers remain unsolved. We aimed to describe the spatial patterns of ants' diversity in the Jamapa River watershed in Mexico based on species distribution models of 253 ant species. Moreover, we intend to understand the relationship between ant diversity with the current climatic conditions and historical climatic stability. The watershed includes maximum elevations of 5636, 4282, and 2500 m from south to north and is exposed to the oceanic northeastern winds. Elevational patterns of species richness were consistent with theoretical predictions but changed south (mid-elevation peaks) to north (monotonic decrease) in the watershed. Species richness was positively associated with mean annual temperature and precipitation seasonality. Climatically stable sites harbored high species richness below 2100 m of elevation but low species richness above 2400 m, while the beta diversity showed the opposite pattern. We observed that present climatic conditions and historical climatic stability were essential drivers of ant diversity in the Jamapa River watershed. Elevational patterns of species richness were concordant with theoretical expectations. They changed from a monotonic decrease of diversity to marked mid-elevation peaks of species richness, presumably associated with the maximum height of the mountains.

Introduction

Despite 220 years of investigation, Humboldt's enigma "What causes tropical mountains to be hyper-diverse?" remains partially unsolved (Rahbek et al., 2019). The juxtaposition of climatic zones within the topography of tropical mountains is considered one of the leading causes of the high biodiversity of tropical mountains (Rahbek et al., 1997; Ohlemüller et al., 2008). Climatic diversity in tropical mountains is associated with mountain ridge exposure to the predominant winds, the height of the peaks, and protected valleys (Rahbek et al., 2019).

Windward slopes of high mountains block oceanic humid winds, causing rain to fall at mid-elevation and potentially favoring net primary productivity (an ecological response) and diversity. On the other hand, inversion layers of cold air can favor stable climates in wind-protected valleys. Such valleys may have served as refuges (museums) and favored diversification (cradles) because of climate stability (evolutionary responses) (Fjeldså, 1999; Rahbek et al., 2019). Further, if valleys remain isolated long enough, high beta diversity among valleys is likely to increase the gamma diversity of tropical mountains (Körner, 2007).



Despite multiple examples of diverse tropical mountains, the contribution of the many factors driving diversity (e.g., peak heights, juxtaposed climates, productivity, museums, cradles, extinctions) needs to be clarified in most cases, limiting our understanding of local biodiverse tropical mountains. Because different mountains have different topographical, geological, and climatic characteristics (Körner et al., 2017), diversity patterns must be compared with caution (c.f., Sanders & Rahbek, 2012). Further, distinct biological groups respond differently to environmental gradients along the mountains (Rahbek, 1995; Graham et al., 2014; Perrigo et al., 2020). Spatial patterns of diversity in tropical mountains include monotonical decreases of diversity with elevation (Grytnes & McCain, 2007) and mid-elevation peaks of diversity (Rahbek & Graves, 2001; Lomolino, 2001), while other studies found no significant changes in diversity from low to relatively high elevations (c.f., Rahbek et al., 2019).

Therefore, research within watersheds with peaks of varied heights may prove advantageous in understanding the diversity of tropical mountains. The Jamapa River watershed in central Veracruz and eastern Puebla states in Mexico combine convenient traits to investigate diversity patterns from the lowlands up to the heights of the mountains. The Jamapa River watershed faces the humid Atlantic Ocean winds, blocked on the west side by mountains that decrease in height from south to north (5636, 4282, and 2500 m asl). Such differences in the height of the mountains are likely to create different environmental gradients at mid-elevation from north to south, with consequent effects on the spatial patterns of the diversity across the mountains of the Jamapa River watershed (c.f., Rahbek & Graves, 2001; Lomolino, 2001).

In this study, we used ecological niche models (Milesi & López, 2005) and species distribution models (Peterson & Soberón, 2012) of ants (Formicidae) to investigate the spatial patterns of diversity in the elevational range of a windward watershed. Ecological niche and species distribution models are robust methods to fill gaps in the Wallacean shortfall of the biological diversity, i.e., the spatial distribution of taxa (Hortal et al., 2015). These models rely on climatic and other abiotic environmental variables to generate hypotheses on the distribution of taxa. Thus, inferring the spatial distribution of a whole taxon based on climatic variables across a complete elevational gradient can help us understand the influence of climate on maintaining the high diversity of tropical mountains.

As a group, ants are found in all land ecosystems but in arctic zones. Ant diversity concentrates in tropics climates (Vásquez, 2011; Schultheiss et al., 2022), but few ant species ascend above 4000 m in elevation (Weber, 1943; Smith, 2015). Roughly half of the studies on ants' diversity along elevational gradients (18 mountain ranges worldwide) reported monotonic decreases in taxonomic diversity. Just over 25% of the studies reported mid-elevation peaks of species richness, whereas other studies reported no significant trends (Smith, 2015).

Ants are taxonomically well-studied in the Jamapa River watershed as specialists in the local research institute (INECOL) and other institutions have conducted research within the watershed for over 30 years (e.g., Rojas-Fernandez, 2001; Valenzuela-González et al., 2013; Ríos-Casanova, 2014; Dátilo et al., 2020; Pérez-Toledo et al., 2022). Therefore, records of species of ants inhabiting the Jamapa River watershed are considered complete. In this study, we aim to fill the Wallacean shortfall of ants in the Jamapa River watershed and investigate the distributional patterns of the alpha and beta diversity concerning current climatic conditions and the historical climatic stability in the watershed. We seek to address the following research questions: (1) Are elevational patterns of ant species richness the same north to south of the Jamapa River watershed? (2) Do elevational patterns of ant species richness correlate with mean temperature and annual precipitation in the Jamapa River watershed? (3) Is beta diversity high at mid-elevations where different climatic regimes and vegetation types juxtapose (ecotones and canyons) in the Jamapa River watershed? (4) Are diversity patterns associated with climatic stability (precipitation or temperature) in the watershed?

Because humid oceanic winds blow predominately from the northeast, we predict that mid-elevation peaks of species richness will be associated with the high peaks (Pico de Orizaba: 5636 m asl, and Cofre de Perote: 4282 m asl) in the south and central section of the Jamapa River watershed, respectively. In contrast, we expect a monotonic decrease of diversity with elevation in the north section of the watershed as mountains are considerably lower than in the central and southern sections. Since elevational changes bring about changes in climate regimes and vegetation types (Hoorn et al., 2018), we expect high beta diversity in juxtaposed climatic zones where transitions of vegetation types happen, ecotones (Carvajal-Hernandez & Krömer, 2015; Gómez-Díaz et al., 2017) and protected valleys and canyons (Peterson et al., 1997). Also, we predict that diversity (alpha and beta) will be high in climatically stable zones of the watershed.

Material and Methods

Study site

The Jamapa River watershed (CNA, 1998; CONABIO, 2001) is in central Veracruz (18.74438 N, 97.26711 W and 19.74529 N, 95.91748 W), covers an area of 10,314 km² (ca. 0.5% of continental Mexico) extending from the windward slopes of the eastern mountains in the Trans Mexican Volcanic Belt: Pico de Orizaba (highest point, 5636 m asl) in the south of the watershed, Cofre de Perote (4282 m asl) in the central part and Astillero (2 m asl) in the north of the watershed. Humid oceanic winds blow predominantly from the northeast, and the mountains form a barrier in the west, very much running north to south (Körner et al., 2011; Salinas-Rodríguez et al., 2022).

Species selection and occurrence records

Based on the data paper of Dátillo et al. (2020), there were 282 ant species in the Jamapa River watershed (Table S1), of which 13 were introduced species and were excluded from the analyses (Table S2). We used all the records in the data paper and the available records in the Global Biodiversity Information Facility for all species native to Mexico reported for the Jamapa River watershed. We discarded all records lacking coordinates, those georeferenced out of the mainland of the American continent and those georeferenced away from the distributional range of each species. Then, we did a geographic filtering to remove duplicated records in the same grid cell ($30''$ arc). For those records collected before or after the time range covered by the climatic variables (1970-2000), we checked if the observed values for bio1, bio12, and bio15 (mean temperature, precipitation, and precipitation seasonality) fell within the interquartile range of all other records of the species. Those records with either of the three climatic variables beyond the interquartile ranges were discarded. After data cleaning, we modeled 253 species verified to be taxonomically resolved (Fisher & Fong, 2023), including 14 species with 1-9 spatially independent records (c.f., Stockwell & Peterson, 2002).

Environmental variables

For the entire American continent, we used the 19 bioclimatic variables from the WorldClim-2 project with a grain of $30''$ arc resolution (Fick & Hijmans, 2017). We used the function `geodata::worldclim_title` in the R programming language (R Core Team, 2022). We downloaded ten tiles to cover the continent. To these climatic tiles, we added layers of soil physical and chemical properties (bulk density, percentage content of slit, clay, sand, and gravel, and total carbon and organic carbon) of the topsoil (0-30 cm) and subsoil (30-100 cm) as soil properties are likely determinants of nesting for many ant species in the tropics (Ryder-Wilkie et al., 2010). Then, we used the `raster::extract` function to obtain each record's climatic variables (19 bioclimatic and 14 soil variables).

Accessibility area

To fulfill the Wallacean shortfall, knowing the entire distributional range of the species, we used a conservative approach in defining the accessibility area of the species, i.e., the historical area at reach through dispersion events for the species and where it is likely to occur, providing the environmental conditions are met (Barve et al., 2011). To this end, we used the Eco-provinces of Canada (Marshall et al., 1999), the Ecological Regions of the United States L-III (Omernik, 1987), the Biogeographic Provinces of Mexico (Morrone et al., 2017), and the Biogeographic Provinces of the Neotropics (Morrone, 2014). Polygons in the border of those regions were merged to eliminate political boundaries. Every polygon with two or more species records was considered part of the accessibility area. Then, based on the accessibility

area, we randomly selected four times as much presence data (c.f., Barbet-Massin et al., 2012) of background points. As described above, we extracted the environmental variables for the background points.

Calibration and evaluation of ecological niche models

We conducted the calibration of the ENMs in two phases. The first phase consisted of applying a machine learning algorithm, random forest, on the presence and background points with 1000 iterations. Each tree processed a random subset of 80% of the presence-background points (response variable) and the 33 environmental variables (predictor variables). Each tree is a computationally exhaustive procedure in which each possible variable and cut-off point is evaluated independently and, therefore, is not affected by the correlation between climatic variables. The process is repeated for each partition of the data set until the internal variation of the subgroups exceeds the difference between their averages. Out of the 1000 trees, we recovered the most likely topology and the environmental variables that separated observation points from background points. Because highly correlated climatic variables will compete in every partition, with only one selected, the resulting consensus tree is unlikely to emerge from highly correlated variables (De'ath & Fabricius, 2000). We coded the procedure in R, which is available in the supplementary material (Supplementary 1).

In the second stage, we used the maxent algorithm to model the species' niche with the set of climatic variables selected by the random forest (Phillips & Dudik, 2008). This algorithm relies on the maximum entropy principle to calculate the most likely distribution of environmental suitability as a function of the difference between the localities of occurrence and the environment of the accessibility area. For the training stage, 60% of the data (presence-background points) was used through the `dismo::maxent` function (Hijmans et al., 2017) of the R statistical software (R Core Team, 2022).

We used the reserved 40% data set to validate the model. We calculated the receiver operating characteristic curve (ROC) and then the partial area under the curve (pAUC), which evaluates the model's ability to discriminate between presences and background points (Peterson et al., 2008). When randomly choosing a pair of cases, this test measures the probability that the model awards a higher probability to presence than to background for all possible pairs in the sample (van Proosdij et al., 2016). Subsequently, to assess the statistical significance of the AUC values, we performed the Mann-Whitney U test using the `Wilcox-test` function of the R statistical software (R Core Team, 2022).

Potential distribution models

For each of the models generated by species that resulted in a statistically significant AUC value ($p < 0.05$), we transformed the continuous suitability projection to a binary projection based on the cut-off threshold of the 10th percentile of the calibration records (Bossò et al., 2013).

Spatial patterns of diversity

Based on the predicted distribution of each species within the watershed, 253 occurrence maps generated in this study were considered to construct a presence/absence matrix (PAM) with 12749 rows (raster cells in the Jamapa River watershed) and 253 columns corresponding to each of the modeled species and an extra column with the ID of the cells. Then, the sum of each row in the matrix, excluding the ID of the cells, was spatially projected to obtain a species richness raster. Also, we calculated beta diversity considering the composition of each focal cell and its 16 neighboring cells. We employed Sørensen's dissimilarity coefficient for this comparison, using the betapart::beta.pair function (Baselga et al., 2018) of the R statistical software (R Core Team, 2022). The Sørensen index allowed us to relate the number of shared species to the arithmetic mean of the species in the focal cell and the neighboring cells.

Climatic stability

The coefficient of variation of the monthly precipitation estimates and mean temperature from three periods. The last interglacial period was around 120,000 years before the present (Otto-Bliesner et al., 2006); the last glacial maximum was 21,000 years in the past, and the mid-Holocene was 6,000 years ago (Braconnot et al., 2007). Then, estimates of climatic stability on precipitation and mean temperature were multiplicatively combined in a single raster.

Statistical analysis

We used generalized additive models (GAM) to test the relationship of species richness with the main climatic variables (mean temperature, annual precipitation, and precipitation seasonality). GAMs are an extension of traditional generalized linear models that incorporate smoothing. We used the mgcv::gam function (Wood & Wood, 2015) in R (R Core Team, 2022). We defined six transects running East to West to account for potential changes in species richness relationships with climatic variables due to large volcanos in the south and central sections of the Jamapa River watershed. We used the transects as a fixed factor in the model. We fitted the model with penalized cubic regression spline smooth that allows the fitting of humps (Wood, 2017). We estimated the parameters under the gamma probability distribution with the inverse link function. For climatic stability, we fitted non-parametric generalized linear models. Models were fitted using the rank transformation of the response variables (alpha and beta diversities) following the logic and algebraic proofs of Conover and Ima (1981) and previous experiences (Ruiz-Guerra et al. 2010; González de León et al. 2016). We used the ID of 11 elevational belts as the primary fixed factor. Elevational belts were 100 m wide and separated at least 100 m from any other belt (0-100, 200-300, 400-500, 1000-1100, 1600-1700, 1800-1900, 2000-2100, 2200-2300, 2400-2500,

2600-2700, 2800-2900). Each belt's sites (raster cells) were classified as low or high climatic stability based on the 0.25 threshold. The analysis did not include belts with less than five sites of either category. The low/high climatic stability factor was nested within the elevational transects to restrict the model to the comparisons in diversity between climatic stability levels within each transect (Sokal & Rolhf, 1987).

Results

Of the 253 modeled species, the species with the smallest predicted distribution in the watershed was *Nylanderia parvula* (< 0.04%), restricted to a few sites between 410 and 460 m altitude in the southern end. In contrast, nine species (ca. 3. 5% of the modeled species) were predicted to occupy at least 95% of the watershed (*Camponotus senex*, *Crematogaster corvina*, *Hypoponera opacior*, *Labidus praedator*, *Neivamyrmex swainsonii*, *Pheidole harrisonfordi*, *Pheidole oaxacana*, *Pseudomyrmex boopis* and *Stenamma felixi*. Also, 149 species (58.8%) would occupy at least 50% of the area in the watershed (Table S3).

Alpha diversity

The geographic pattern of predicted richness in the Jamapa River watershed showed a gradient in which richness decreased from the lowlands (east) to the highlands (west) but also revealed a marked increase in potential richness at mid-elevations, between 600 and 1400 m elevation in the southern section of the watershed, and between 800 and 1400 m elevation in the central section, while in the northern section, species richness gradually decreased from the lowlands to the highlands (Figure 1).

The GAM models showed a positive correlation between mean annual temperature (BIO1) and potential species richness with R^2 -adjusted = 0.90. Above 15 °C, all six transects showed mid-range peaks in species richness, with peaks increasing in magnitude from north to south of the watershed, and the highest points of the peaks were between 17 and 21 °C (Figure 2a). While precipitation (BIO12) showed no correlation with species richness (Figure 2b), species richness increased as precipitation seasonality (BIO15) also increased and was similar in the five analyzed transects (Figure 2c).

Species richness also showed a mixed pattern regarding climatic stability (Figure 3). The lowest-elevation belt (0-100 m) showed no significant differences between low and high climatic stability sites (Figure 3a). In contrast, the remaining elevation segments below 2100 m showed significant differences between low and high-climatic stability areas. Areas of high climatic stability harbored more species richness than those of low climatic stability (Fig. 3b-g). On the other hand, elevational segments above 2300 m showed the opposite pattern with low species richness in areas of high climatic stability compared to low climatic stability areas where higher species richness was predicted (Fig 3i-k).

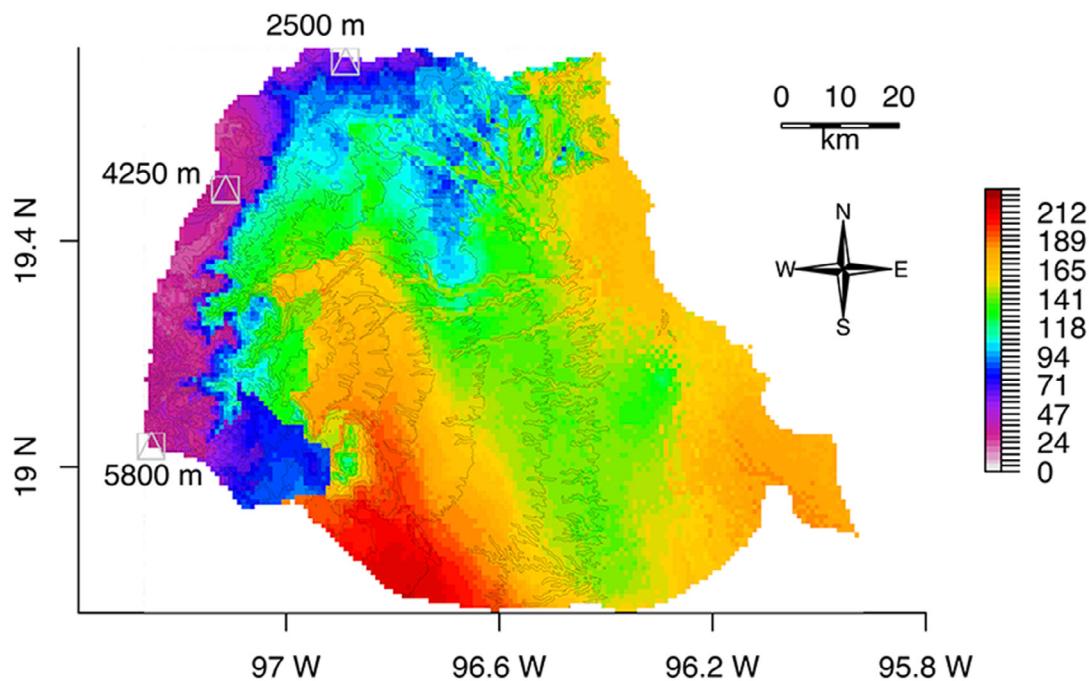


Fig 1. Predicted potential ant species richness in the Jamapa River watershed. Elevation intervals are separated by 200 m.

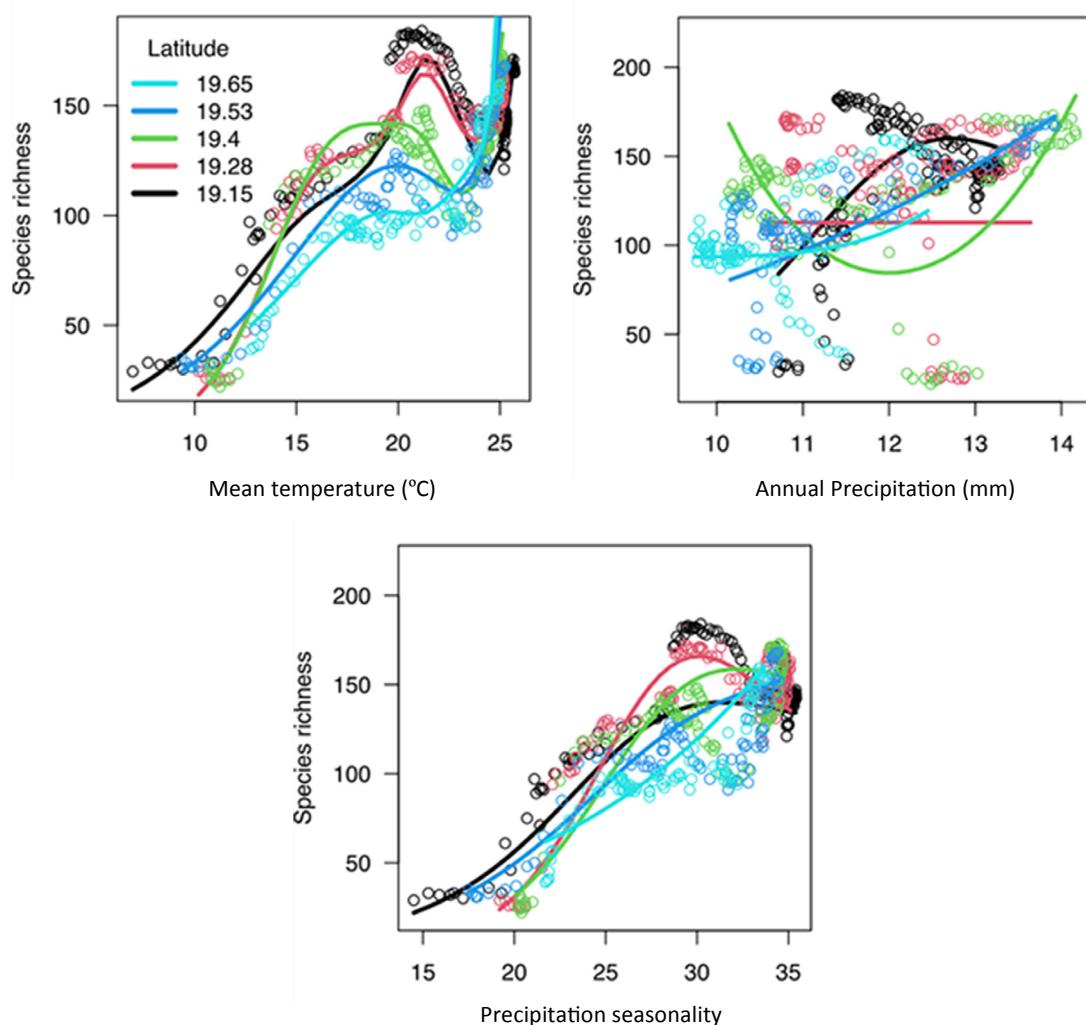


Fig 2. Relationship of ant species richness with (a) mean annual temperature (BIO1) with $r^2 = 0.93$, (b) annual precipitation (BIO12) with $r^2 = 0.13$ and (c) precipitation seasonality (BIO15) with $r^2 = 0.85$, using five elevational transects at different latitudes. Lines were fitted with generalized additive models based on logarithmic values of species richness and predictions back transformed for graphical representation.

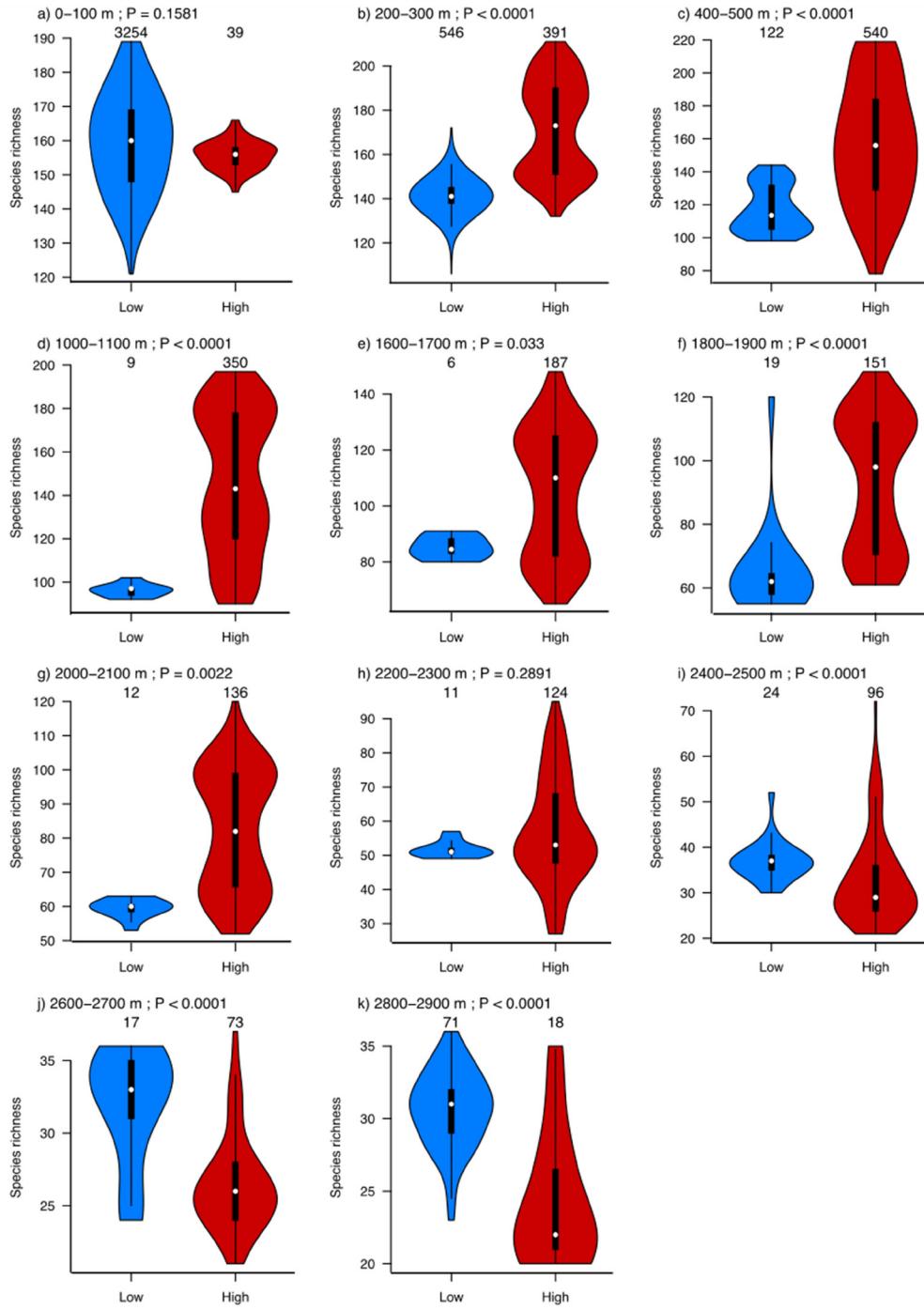


Fig 3. Ant species richness as a function of climatic stability across 11 elevational 100 m wide belts (a-k). Low climatic stability was defined as < 0.25 , while high climatic stability was > 0.25 . probability values were derived from contrast tests (a t-test in which the standard error is estimated from the linear predictor of the model) between the two climatic stability categories in each belt. The model was a generalized linear model with a gamma error structure and inverse link function.

Beta diversity

The beta diversity of the cells about their 16 neighboring cells was low (Figure 4). Fifty-four percent of the area in the Jamapa River watershed had a Sørensen dissimilarity coefficient value less than 0.025, and only 10.2% of the cells in the watershed had a value greater than 0.1. High beta diversity was observed in the upper part of the elevation gradient, the north of the watershed (0.1 to 0.2), and some gorges in the central part (0.05 to 0.1).

Regarding climatic stability, we observed that at the lowest elevation belt (0–100 m), beta-diversity was higher in high climatically stable sites (Figure 5a). However, in the elevation belts between 200 and 1900 m, we observed higher beta-diversity in less climatically stable sites compared to climatically stable sites (Figure 5b-f). Then, from 2200 up to 2900 m, more climatically stable sites were, again, more beta diverse than less climatically stable zones (Figure 5h-k).

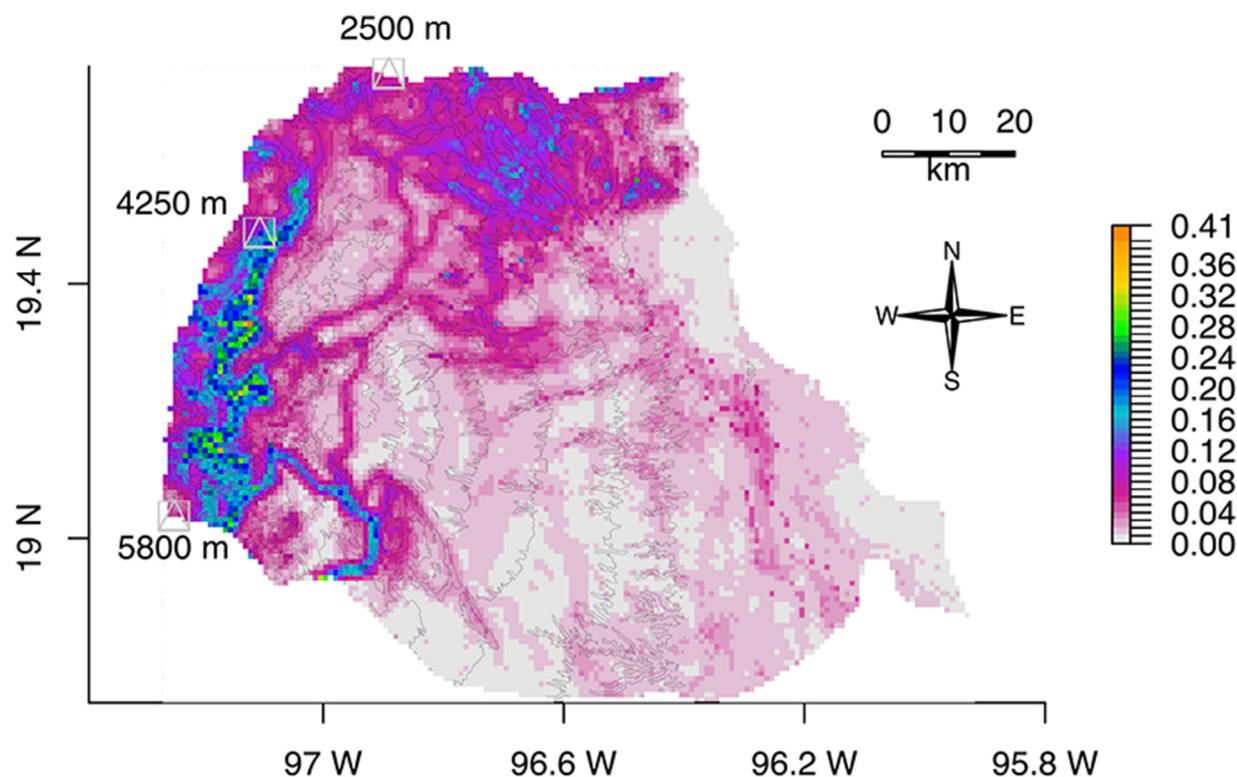


Fig 4. Prediction of Sørensen beta diversity of ants in the Jamapa River watershed. Elevation intervals are separated by 200 m.

Discussion

The Jamapa River watershed includes full elevational gradients from sea levels up to 2500 and 4658 m and a wide range of biodiversity distributional patterns across the lowlands and the mountains. The northern portion of the watershed showed the most straightforward expected pattern for species richness. Here, species richness decreased monotonically from the lowlands to the mountains at most 2500 m in elevation. Further, species richness correlated linearly and positively with mean temperature. This elevational pattern has been reported for many taxa, including lake sediment bacteria (Zeng et al., 2016), woody vegetation (Cirimwami et al., 2019; Griffiths et al., 2021), reptiles (McCain, 2010) and ants (Kunene et al., 2022), although what are the actual drivers of changes in species richness still unclear and likely vary among taxonomic and functional groups, the specific location and orientation of the watersheds in the mountains (Kluge et al., 2006; Irl et al., 2015; Song et al., 2021). In this study, mean temperature correlated significantly with ant species richness, while precipitation did not.

In the central and southern sections of the watershed, where mountains reached well above 4000 m, there were evident mid-elevation peaks in species richness, especially in the southern end of the watershed, where species richness halfway up the mountains was higher than anywhere else. Mid-elevation humps of diversity have been observed in other tropical mountain, including various biological groups

such as vascular epiphytes (Krömer et al., 2005; Cardelús et al., 2006), vascular plants (Zhou et al., 2019), ferns (Syfert et al., 2018) and also ants (Nowrouzi et al., 2016). In our study, mid-elevation peaks in species richness also correlated closely with mean temperature, although trends were nonlinear. Mark humps in species richness were observed between 17 and 21°C, with humps occurring at warmer temperatures towards the south of the watershed. This pattern suggests that mountain height may push mid-elevation peaks of diversity down the mountain to warmer conditions. Besides, changes in temperature with elevation, high mountains exposed to humid winds from the ocean block clouds and cause high precipitation at mid-elevations, which has been taught as one of the drivers of high productivity and diversity (Currie, 1991; Kreft & Jetz, 2007; Rapp & Silman, 2012).

Further, species richness increased with precipitation seasonality, suggesting different phenologies among ant species could contribute to increased species richness. Ants's phenology may respond to the availability of different resources during the seasons in highly seasonal sites (Andersen, 1986). This pattern is concordant with the known diversity of ants not only in terms of species but also in diets and behavior that allow ants to distribute in most terrestrial environments (Decaëns et al., 2002; Lach et al., 2010; Del Toro et al., 2012;). This pattern has also been observed for other biological groups, such as lianas, which are more species-rich in seasonally dry forests than evergreen forests in Panama (Parolari et al., 2020).

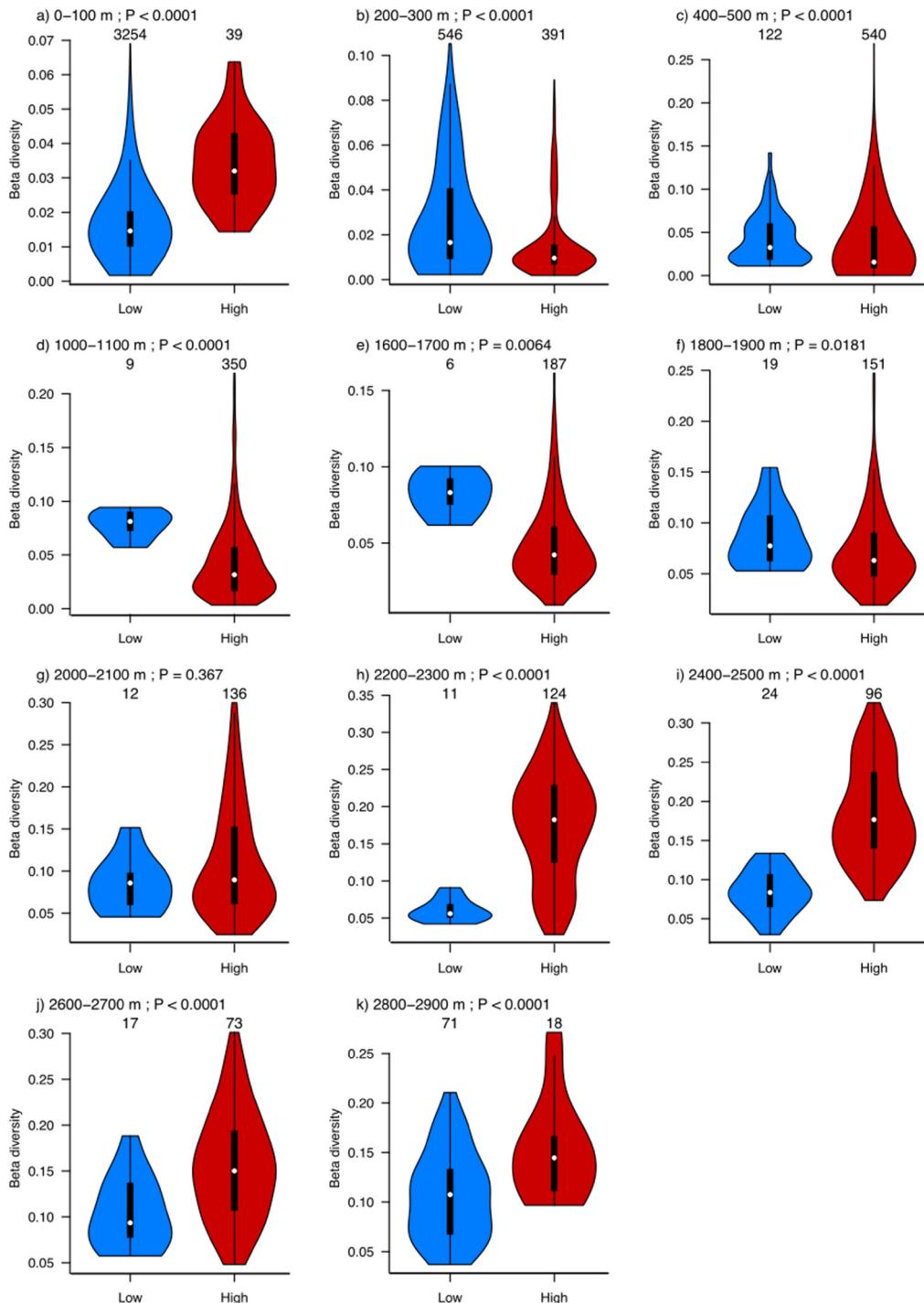


Fig 5. Sørensen beta diversity of ants as a function of climatic stability across 11 elevational 100 m wide belts (a-k). Low climatic stability was defined as < 0.25 , while high climatic stability was > 0.25 . probability values were derived from contrast tests (a t-test in which the standard error is estimated from the linear predictor of the model) between the two climatic stability categories in each belt. The model was a generalized linear model with a gamma error structure and inverse link function.

Alternatively, long-lasting climatic stability favors niche specialization along with low extinction rates, which may vary among lineages, thus favoring local alpha-diversity with a blend of derive and relic taxa (Werneck et al., 2012; Gao & Liu, 2018), although there is some evidence showing than the distribution of derived and old and relic lineages are spatially separated (Sonne et al. 2022; González-Orozco 2023; Gopal et al. 2023). Nonetheless, cloud forests, located halfway up the mountains, are species-rich zones and harbor numerous

species of restricted distribution (Cooper et al. 2022; Sonne et al. 2022; González-Orozco 2023). Here, we evaluated the climatic stability over the past 120,000 years and found it correlates with alpha and beta-diversity, but trends changed from the low land to the heights in the mountains. Species richness was high in climatically stable sites in the lowlands and up to 2100 m in the mountains. Above 2300 m of elevation, climatically stable zones harbored low species-rich communities than more climatically variable zones.

Regarding beta diversity, we observed the opposite pattern: low beta diversity in climatically stable zones in the lowlands and up to 2000 m in elevation and higher beta diversity in climatically stable zones above 2200 m in elevation. This finding suggests that at 2100 m of elevation and below, climatically stable zones accumulate a high number of species, presumably because of niche specialization and the conservation of old lineages favored by predictable climatic conditions, while above 2300 m of elevation, climatically stable zones harbored fewer species than those zones with more considerable climatic variability. This contradictory result suggests that climatic stability above 2300 m means constantly unfavorable climatic conditions that may limit resource availability (Girardin et al., 2010) and, therefore, low species richness. In contrast, climatically variable zones high in the mountains may allow species to coexist with different phenologies or a relatively large number of species to use resources during the more favorable season (Mellard et al., 2019). Both possibilities need further investigation.

The use of predictive techniques to explore geographical patterns of the distribution of diversity is a valuable opportunity to fulfill the Wallacean shortfall, i.e., on the distribution of biodiversity. Evidence supports the congruency of patterns based on actual records, and those based on species distribution models are well correlated (Sosa & Loera, 2017). However, alpha diversity richness may be overestimated (Mateo et al., 2012). Nonetheless, using predictive models of the distribution of ants in the Jamapa River watershed revealed spatial patterns of alfa beta diversity that match the predictions and further highlight the influence of high mountains in producing diversity peaks at mid-elevations. In the south of the watershed, the mid-elevation peaks of diversity occurred at warmer temperatures than those in the central section (south to north) of the watershed.

In contrast, in the north of the watershed, there were no mid-elevation peaks of alpha diversity. Thus, elevational patterns of diversity are not only susceptible to the specific geographic location of the mountains (Sanders & Rahbek, 2012; Guo et al., 2013; Körner et al., 2017) but also the specifics of the taxonomic or functional group surveyed (Rahbek, 1995) but also the specific location and orientation of elevational transects within the mountains or whole watersheds. Therefore, achieving a more comprehensive understanding of the spatial patterns of biodiversity across elevational gradients requires multiple transects in each region to adequately represent the heterogeneity that the height of mountains may cause.

Overall, we showed that niche and species distribution models of the myrmecofauna of the Jamapa River watershed produced elevational patterns of species concordant with theoretical expectations. Further, we showed that those patterns change from a monotonic decrease in diversity to marked mid-elevation peaks of species richness, presumably associated with the maximum high of the mountains that intercept the predominantly northeastern winds from the Gulf of Mexico. Further, we observed that present and historical

climatic conditions are essential drivers of ants diversity in the Jamapa River watershed.

Authors' Contributions

SP-B: investigation; resources; visualization; writing-original draft preparation; writing-review and editing.

AL-N: conceptualization; methodology; resources; supervision; writing-review and editing.

RG: conceptualization; data curation; formal analysis; investigation; methodology; resources; software; supervision; validation; visualization; writing-review and editing.

Acknowledges

We thank two anonymous reviewers for their comments on an early draft of the manuscript. This investigation was funded by INECOL and the AWE's benefactor program on behalf of the first author of the manuscript.

References

- Andersen, A.N. (1986). Diversity, seasonality, and community organization of ants at adjacent heath and woodland sites in southeastern Australia. *Australian Journal of Zoology*, 34: 53-64.
- Barbet-Massin, M., Jiguet, F., Albert, C.H., & Thuiller, W. (2012). Selecting pseudo-absences for species distribution models: How, where and how many? *Methods in ecology and Evolution*, 3: 327-338.
- Barve, N., Barve, V., Jiménez-Valverde, A., Lira-Noriega, A., Maher, S.P., Peterson, A.T., Soberón, J. & Villalobos, F. (2011). The crucial role of the accessible area in ecological niche modeling and species distribution modeling. *Ecological Modelling*, 222: 1810-1819.
- Baselga, A., Orme, D., Villeger, S., De Bortoli, J., Leprieur, F. & Baselga, M.A. (2018). Package 'betapart'. Partitioning beta diversity into turnover and nestedness components, version, 1.
- Bosso, L., Rebelo, H., Garonna, A.P. & Russo, D. (2013). Modelling geographic distribution and detecting conservation gaps in Italy for the threatened beetle *Rosalia alpina*. *Journal for Nature Conservation*, 21: 72-80.
- Braconnot, P., Otto-Bliesner, B., Harrison, S., Joussaume, S., Peterchmitt, J. Y., Abe-Ouchi, A., ... & Zhao, Y. (2007). Results of PMIP2 coupled simulations of the Mid-Holocene and Last Glacial Maximum-Part 1: experiments and large-scale features. *Climate of the Past*, 3: 261-277.
- Cardelús, C.L., Colwell, R.K. & Watkins Jr, J.E. (2006). Vascular epiphyte distribution patterns: explaining the mid-elevation richness peak. *Journal of Ecology*, 144-156.
- Carvajal-Hernández, C.I. & Krömer, T. (2015). Riqueza y distribución de helechos y licófitos en el gradiente altitudinal del Cofre de Perote, centro de Veracruz, México. *Botanical Sciences*, 93: 601-614.

- Cirimwami, L., Doumenge, C., Kahindo, J. M. & Amani, C. (2019). The effect of elevation on species richness in tropical forests depends on the considered lifeform: results from an East African mountain forest. *Tropical Ecology*, 60: 473-484.
- Comisión Nacional del Agua (CNA), (1998). Cuencas Hidrográficas [vector file] 1:250000. México. https://gisviewer.semarnat.gob.mx/aplicaciones/Atlas2015/agua_chidrograficas.html
- Comisión Nacional para el Conocimiento y Uso de la Biodiversidad (CONABIO), (2001). Cuencas Hidrológicas de la República Mexicana [vector file] 1:250000. México. http://www.conabio.gob.mx/informacion/gis/?vns=gis_root/hidro/chidro/cue250kgw.
- Conover, W.J. & Iman, R.L. (1981). Rank transformations as a bridge between parametric and nonparametric statistics. *American Statistician*, 124-129.
- Cooper, J.C., Crouch, N.M., Ferguson, A.W. & Bates, J.M. (2022). Climatic refugia and reduced extinction correlate with underdispersion in mammals and birds in Africa. *Ecology and Evolution*, 12: e8752.
- Currie, D.J. (1991). Energy and large-scale patterns of animal- and plant-species richness. *The American Naturalist*, 137: 27-49.
- Dátillo, W., Vásquez-Bolaños, M., Ahuatzin, D.A., Antoniazzi, R., Chávez-González, E., Corro, E., Luna, P., Guevara, R., Villalobos, F., Madrigal-Chavero, R. and Falcão, J.C.D.F. (2020). Mexico ants: incidence and abundance along the Nearctic-Neotropical interface. *Ecology*, 101: e02944.
- Decaëns, T., Asakawa, N., Galvis, J.H., Thomas, R.J. and Amézquita, E. (2002). Surface activity of soil ecosystem engineers and soil structure in contrasted land use systems of Colombia. *European Journal of Soil Biology*, 38: 267-271.
- Del Toro, I., Ribbons, R.R. & Pelini, S.L. (2012). The little things that run the world revisited: a review of ant-mediated ecosystem services and disservices (Hymenoptera: Formicidae). *Myrmecological News*, 17: 133-46.
- De'ath, G. & Fabricius, K.E. (2000). Classification and regression trees: a powerful yet simple technique for ecological data analysis. *Ecology*, 81: 3178-3192.
- Fick, S.E. & Hijmans, R.J. (2017). WorldClim 2: new 1 km spatial resolution climate surfaces for global land areas. *International Journal of Climatology*, 37: 4302-4315.
- Fisher, B. & Fong, J. (2023). AntWeb. California Academy of Sciences. Occurrence dataset. <https://doi.org/10.15468/wqmjjt>
- Fjeldså, J., Lambin, E. & Mertens, B. (1999). Correlation between endemism and local ecoclimatic stability documented by comparing Andean bird distributions and remotely sensed land surface data. *Ecography*, 22: 63-78.
- Gao, J. & Liu, Y. (2018). Climate stability is more important than water-energy variables in shaping the elevational variation in species richness. *Ecology and Evolution*, 8: 6872-6879.
- Girardin, C.A.J., Malhi, Y., Aragão, L.E.O.C., Mamani, M., Huaraca Huasco, W., Durand, L., Feeley, K.J., Rapp, J., Silva-Espejó, J.E., Silman, M., Salinas, R. & Whittaker, R. J. (2010). Net primary productivity allocation and cycling of carbon along a tropical forest elevational transect in the Peruvian Andes. *Global Change Biology*, 16: 3176-3192.
- Gómez-Díaz, J.A., Krömer, T., Kreft, H., Gerold, G., Carvajal-Hernández, C.I. & Heitkamp, F. (2017). Diversity and composition of herbaceous angiosperms along gradients of elevation and forest-use intensity. *PloS One*, 12: e0182893.
- González de Leon, S., Herrera, I. & Guevara, R. (2016). Mating system, population growth, and management scenario for Kalanchoe pinnata in an invaded seasonally dry tropical forest. *Ecology and Evolution*, 6: 4541-4550.
- González-Orozco, C.E. (2023). Unveiling evolutionary cradles and museums of flowering plants in a neotropical biodiversity hotspot. *Royal Society Open Science*, 10: 230917.
- Gopal, A., Bharti, D. K., Page, N., Dexter, K. G., Krishnamani, R., Kumar, A. & Joshi, J. (2023). Range restricted old and young lineages show the southern Western Ghats to be both a museum and a cradle of diversity for woody plants. *Proceedings of the Royal Society B*, 290: 20222513.
- Graham, C. H., Carnaval, A. C., Cadena, C. D., Zamudio, K. R., Roberts, T. E., Parra, J. L., McCain, C.M., Bowie R.C.K., Moritz, C., Baines, S.B., Schneider, J., VanDerWal, J., Rahbek, C., Kosak, K.H. & Sanders, N. J. (2014). The origin and maintenance of montane diversity: integrating evolutionary and ecological processes. *Ecography*, 37: 711-719.
- Griffiths, A.R., Silman, M.R., Farfan-Rios, W., Feeley, K.J., Cabrera, K.G., Meir, P., Salinas, N., Segovia, R.A. & Dexter, K.G. (2021). Evolutionary diversity peaks at mid-elevations along an Amazon-to-Andes elevation gradient. *Frontiers in Ecology and Evolution*, 9: 680041.
- Grytnes, J.A. & McCain, C.M. (2007). Elevational trends in biodiversity. *Encyclopedia of biodiversity*, 2, pp.1-8.
- Guo, Q., Kelt, D. A., Sun, Z., Liu, H., Hu, L., Ren, H. & Wen, J. (2013). Global variation in elevational diversity patterns. *Scientific Reports*, 3: 3007.
- Hijmans, R.J., Phillips, S., Leathwick, J., Elith, J. & Hijmans, M.R.J. (2017). Package 'dismo'. Circles, 9: 1-68.
- Hoorn, C., Perrigo, A. & Antonelli, A. (2018). Mountains, climate and biodiversity. John Wiley & Sons, UK.
- Hortal, J., de Bello, F., Diniz-Filho, J.A.F., Lewinsohn, T.M., Lobo, J.M. & Ladle, R.J. (2015). Seven shortfalls that beset large-scale knowledge of biodiversity. *Annual Review of Ecology, Evolution, and Systematics*, 46: 523-549.
- Irl, S.D., Harter, D.E., Steinbauer, M.J., Gallego Puyol, D., Fernández-Palacios, J.M., Jentsch, A. & Beierkuhnlein, C. (2015). Climate vs. topography-spatial patterns of plant species

- diversity and endemism on a high-elevation island. *Journal of Ecology*, 103: 1621-1633.
- Lach, L., Parr, C. & Abbott, K. (2010). *Ant ecology*. Oxford university press.
- Kluge, J., Kessler, M. & Dunn, R.R. (2006). What drives elevational patterns of diversity? A test of geometric constraints, climate and species pool effects for pteridophytes on an elevational gradient in Costa Rica. *Global Ecology and Biogeography*, 15: 358-371.
- Körner, C. (2007). The use of “altitude” in ecological research. *Trends in Ecology and Evolution*, 22: 569-574.
- Körner, C., Paulsen, J. & Spehn, E. M. (2011). A definition of mountains and their bioclimatic belts for global comparisons of biodiversity data. *Alpine Botany*, 121: 73-78.
- Körner, C., Jetz, W., Paulsen, J., Payne, D., Rudmann-Maurer, K. & M. Spehn, E. (2017). A global inventory of mountains for bio-geographical applications. *Alpine Botany*, 127: 1-15.
- Kreft, H. & Jetz, W. (2007). Global patterns and determinants of vascular plant diversity. *Proceedings of the National Academy of Sciences*, 104: 5925-5930.
- Krömer, T., Kessler, M., Robbert Gradstein, S. & Acebey, A. (2005). Diversity patterns of vascular epiphytes along an elevational gradient in the Andes. *Journal of Biogeography*, 32: 1799-1809.
- Kunene, C., Foord, S.H., Scharff, N., Pape, T., Malumbres-Olarte, J. & Munyai, T.C. (2022). Ant diversity declines with increasing elevation along the Udzungwa Mountains, Tanzania. *Diversity*, 14: 260.
- Lomolino, M.V. (2001). Elevation gradients of species-density: historical and prospective views. *Global Ecology and Biogeography*, 10: 3-13.
- Marshall, I.B., Schut, P.H. & Ballard, M. (1999). A National Ecological Framework for Canada: Attribute Data. Agriculture and Agri-Food Canada, Research Branch, Centre for Land and Biological Resources Research and Environment Canada, State of the Environment Directorate, Ecozone Analysis Branch. Ottawa/Hull.
- Mateo, R.G., Felicísimo, Á.M., Pottier, J., Guisan, A. & Muñoz, J. (2012). Do stacked species distribution models reflect altitudinal diversity patterns? *PLoS One*, 7: e32586.
- McCain, C.M. (2010). Global analysis of reptile elevational diversity. *Global Ecology and Biogeography*, 19: 541-553.
- Millard, J.P., Audoye, P. & Loreau, M. (2019). Seasonal patterns in species diversity across biomes. *Ecology*, 100: e02627.
- Milesi, F.A. & Lopez de Casenave, J. (2005). El concepto de nicho en ecología aplicada: del nicho al hecho hay mucho trecho. *Ecología Austral*, 15: 131-148.
- Morrone, J.J. (2014). Biogeographical regionalisation of the Neotropical region. *Zootaxa*, 3782: 1-110.
- Morrone, J.J., Escalante, T. & Rodriguez-Tapia, G. (2017). Mexican biogeographic provinces: Map and shapefiles. *Zootaxa*, 4277: 277-279.
- Nowrouzi, S., Andersen, A.N., Macfadyen, S., Staunton, K.M., VanDerWal, J. & Robson, S.K. (2016). Ant diversity and distribution along elevation gradients in the Australian wet tropics: the importance of seasonal moisture stability. *PloS One*, 11: e0153420.
- Ohlemüller, R., Anderson, B.J., Araujo, M.B., Butchart, S.H., Kudrna, O., Ridgely, R.S. & Thomas, C.D. (2008). The coincidence of climatic and species rarity: high risk to small-range species from climate change. *Biology Letters*, 4: 568-572.
- Omernik, J.M. (1987). Ecoregions of the conterminous United States. Map (scale 1:7,500,000). *Annals of the Association of American Geographers*, 77: 118-125.
- Otto-Bliesner, B.L., Marshall, S.J., Overpeck, J.T., Miller, G.H., Hu, A. & CAPE Last Interglacial Project members. (2006). Simulating Arctic climate warmth and icefield retreat in the last interglaciation. *Science*, 311: 1751-1753.
- Parolari, A.J., Paul, K., Griffing, A., Condit, R., Perez, R., Aguilar, S. & Schnitzer, S.A. (2020). Liana abundance and diversity increase with rainfall seasonality along a precipitation gradient in Panama. *Ecography*, 43: 25-33.
- Pérez-Toledo, G.R., Villalobos, F., Silva, R.R., Moreno, C.E., Pie, M.R. & Valenzuela-González, J.E. (2022). Alpha and beta phylogenetic diversities jointly reveal ant community assembly mechanisms along a tropical elevational gradient. *Scientific Reports*, 12: 7728.
- Perrigo, A., Hoorn, C. & Antonelli, A. (2020). Why mountains matter for biodiversity. *Journal of Biogeography*, 47: 315-325.
- Peterson, D.L., Schreiner, E.G. & Buckingham, N.M. (1997). Gradients, vegetation and climate: spatial and temporal dynamics in the Olympic Mountains, USA. *Global ecology and Biogeography Letters*, 6: 7-17.
- Peterson, A.T., Papes, M. & Soberón, J. (2008). Rethinking receiver operating characteristic analysis applications in ecological niche modeling. *Ecological Modelling*, 213: 63-72.
- Peterson, A.T. & Soberón, J. (2012). Species distribution modeling and ecological niche modeling: getting the concepts right. *Natureza e Conservação*, 10: 102-107.
- Phillips, S.J. & Dudík, M. (2008). Modeling of species distributions with Maxent: new extensions and a comprehensive evaluation. *Ecography*, 31: 161-175.
- R Core Team (2023). R: A Language and Environment for Statistical Computing. R Foundation for Statistical Computing, Vienna, Austria. <https://www.R-project.org/>
- Rahbek, C. (1995). The elevational gradient of species richness: a uniform pattern? *Ecography*, pp. 200-205.

- Rahbek, C. (1997). The relationship among area, elevation, and regional species richness in neotropical birds. *The American Naturalist*, 149: 875-902.
- Rahbek, C. & Graves, G.R. (2001). Multiscale assessment of patterns of avian species richness. *Proceedings of the National Academy of Sciences*, 98: 4534-4539.
- Rahbek, C., Borregaard, M.K., Colwell, R.K., Dalsgaard, B.O., Holt, B.G., Morueta-Holme, N., Nogues-Bravo, D., Whittaker, R.J. & Fjeldså, J. (2019). Humboldt's enigma: What causes global patterns of mountain biodiversity? *Science*, 365: 1108-1113.
- Rapp, J.M., & Silman, M.R. (2012). Diurnal, seasonal, and altitudinal trends in microclimate across a tropical montane cloud forest. *Climate Research*, 55: 17-32.
- Ríos-Casanova, L. (2014). Biodiversity of ants in Mexico. *Revista Mexicana de Biodiversidad*, 85: S392-S398.
- Rojas-Fernández, P. (2001). Las hormigas del suelo en México: diversidad, distribución e importancia (Hymenoptera: Formicidae). *Acta Zoológica Mexicana*, 1: 189-238.
- Ruiz-Guerra, B., Guevara, R., Mariano, N.A. & Dirzo, R. (2010). Insect herbivory declines with forest fragmentation and covaries with plant regeneration mode: evidence from a Mexican tropical rain forest. *Oikos*, 119: 317-325.
- Ryder-Wilkie, K.T., Mertl, A.L. & Traniello, J.F. (2010). Species diversity and distribution patterns of the ants of Amazonian Ecuador. *Plos One*, 5: e13146.
- Salinas-Rodríguez, M.M., Hernández-Sandoval, L., Carrillo-Reyes, P., Castillo-Gómez, H.A., Castro-Castro, A., Estrada-Castillón, E., Figueroa-Martínez, D.S., Gómez-Escamilla, I.N., González-Elizondo, M., Gutiérrez-Ortega, J.S., Hernández-Rendón, J., Munguía-Lino, G., De-Nova, J.A., Ortíz-Brunel, J.P., Rubio-Méndez, G., Ruíz-Sánchez, C., Sandoval-Mata, T.N., Soltero-Quintana, R., Steinmann, V., Valencia-A, S. & Zamudio-Ruiz, S. (2022). Diversidad de plantas vasculares de la provincia fisiográfica de la Sierra Madre Oriental, México. *Botanical Sciences*, 100: 469-492.
- Sanders, N.J. & Rahbek, C. (2012). The patterns and causes of elevational diversity gradients. *Ecography*, 35: 1.
- Schultheiss, P., Nooten, S.S., Wang, R., Wong, M.K., Brassard, F. & Guénard, B. (2022). The abundance, biomass, and distribution of ants on Earth. *Proceedings of the National Academy of Sciences*, 119: e2201550119.
- Smith, M.A. (2015). Ants, elevation, phylogenetic diversity and community structure. *Ecosphere*, 6: 1-17.
- Sokal, R.R. & Rohlf, F.J. (1987). Biostatistics. Francise & Co, New York, 10.
- Sonne, J., Dalsgaard, B., Borregaard, M.K., Kennedy, J., Fjeldså, J. & Rahbek, C. (2022). Biodiversity cradles and museums segregating within hotspots of endemism. *Proceedings of the Royal Society B*, 289: 20221102.
- Song, X., Cao, M., Li, J., Kitching, R.L., Nakamura, A., Laidlaw, M.J., Tang, Y., Sun, Z., Zhang, W. & Yang, J. (2021). Different environmental factors drive tree species diversity along elevation gradients in three climatic zones in Yunnan, southern China. *Plant Diversity*, 43: 433-443.
- Sosa, V. & Loera, I. (2017). Influence of current climate, historical climate stability and topography on species richness and endemism in Mesoamerican geophyte plants. *PeerJ*, 5: e3932.
- Syfert, M.M., Brummitt, N.A., Coomes, D.A., Bystrakova, N. & Smith, M.J. (2018). Inferring diversity patterns along an elevation gradient from stacked SDMs: A case study on Mesoamerican ferns. *Global Ecology and Conservation*, 16: e00433.
- Stockwell, D.R. & Peterson, A.T. (2002). Effects of sample size on accuracy of species distribution models. *Ecological Modelling*, 148: 1-13.
- van Proosdij, A.S., Sosef, M.S., Wieringa, J.J. & Raes, N. (2016). Minimum required number of specimen records to develop accurate species distribution models. *Ecography*, 39: 542-552.
- Valenzuela-González, J., Parra-Cabañas, A.V., Quiroz-Robledo, L., Martínez-Tlapa, D.L. & Montes-de-Oca-Torres, E.D. (2013). Variación de la mirmecofauna en un gradiente altitudinal en la región central de Veracruz, México. *Formicidae de México* (ed. by Vásquez-Bolaños, M., Castaño-Meneses, G., Cisneros-Caballero, A., Quiroz-Rocha, G.A., Navarrete-Heredia, J.L.), pp. 75-82. Cuerpo Académico de Zoología UDG-CA-51. Guadalajara, Jalisco, México.
- Vásquez-Bolaños, M. (2011). Lista de especies de hormigas (Hymenoptera: Formicidae) para México. *Dugesiana*, 18: 95-133.
- Weber, N.A. (1943). The ants of the Imatong Mountains, Anglo-Egyptian Sudan. *Bulletin of Museum of Comparative Zoology*, 93: 263-289.
- Werneck, F.P., Nogueira, C., Colli, G.R., Sites Jr, J.W. & Costa, G.C. (2012). Climatic stability in the Brazilian Cerrado: implications for biogeographical connections of South American savannas, species richness and conservation in a biodiversity hotspot. *Journal of Biogeography*, 39: 1695-1706.
- Wood, S.N. (2017). Generalized additive models: an introduction with R (2. edition). Chapman and Hall/CRC Press.
- Wood, S. & Wood, M.S. (2015). Package ‘mgcv’. R package version, 1: 729.
- Zeng, J., Zhao, D., Li, H., Huang, R., Wang, J. & Wu, Q. L. (2016). A monotonically declining elevational pattern of bacterial diversity in freshwater lake sediments. *Environmental Microbiology*, 18: 5175-5186.
- Zhou, Y., Ochola, A.C., Njogu, A.W., Boru, B.H., Mwachala, G., Hu, G., Xin, H. & Wang, Q. (2019). The species richness pattern of vascular plants along a tropical elevational gradient and the test of elevational Rapoport's rule depend on different life-forms and phytogeographic affinities. *Ecology and Evolution*, 9: 4495-4503.

Supplementary Tables

Table S1. List of modeled species of ants of the Jamapa River watershed in Central Veracruz, Mexico. 10P, tenth percentile of suitability predictions used as binarization threshold; TSS, true skill statistic; FOR, false omission rate; AUCp, partial area under the curve, including the minimum (Min), first (Q1), second (Q2) and third quartile (Q3) and the maximum value based on 500 random subsets of validation data; P, probability of the one sample Wilcoxon test on AUCp values. The following species also reported for the Jamapa River watershed were excluded from the analyses because of insufficient records: *Adelomyrmex doreae*, *Camponotus conspicuus sharpi*, *Colobopsis abdita*, *C. etiolata*, *C. mathildeae*, *Leptothorax striatulus*, *Dorymyrmex smithi*, *Nylanderia austrococcidua*, *Paratrechina austrococcidua*, *Proceratium convexiceps*, *Strumigenys crenulata*, *S. oconitrilloae*, *S. sevestra*, *S. turpis*, *Temnothorax striatulus* and *Trachymyrmex squamulifer*.

Subfamily/Species	10P	TSS	FOR	AUCp					P
				Min	Q1	Q2	Q3	Max	
Amblyoponinae									
<i>Fulakora orizabana</i> (Brown, 1960)	0.325	0.822	0.022	4.8	5.3	5.7	81.2	437.1	< 0.001
Dolichoderinae									
<i>Azteca alfari</i> Emery, 1893	0.456	0.858	0.012	27.6	39.6	43.5	56.3	228.0	< 0.001
<i>Azteca gnava</i> Forel, 1906	0.373	0.842	0.008	53.0	180.7	352.6	580.8	8321.1	< 0.001
<i>Azteca nigricans</i> Forel, 1899	0.580	0.908	0.008	191.4	673.3	1078.9	2398.7	5038.3	< 0.001
<i>Azteca velox</i> Forel, 1899	0.468	0.811	0.014	34.0	42.8	50.2	61.1	141.3	< 0.001
<i>Dolichoderus diversus</i> Emery, 1894	0.431	0.823	0.024	3.1	3.2	3.4	67.0	226.1	< 0.001
<i>Dolichoderus lutosus</i> (Smith, 1858)	0.394	0.804	0.021	23.9	31.6	37.3	59.9	159.1	< 0.001
<i>Dorymyrmex bicolor</i> Wheeler, 1906	0.242	0.854	0.020	6.8	7.1	7.3	7.5	29.6	< 0.001
<i>Dorymyrmex insanus</i> (Buckley, 1866)	0.448	0.809	0.027	5.3	5.7	5.8	6.6	7.4	< 0.001
<i>Forelius damiani</i> Guerrero & Fernández, 2008	0.204	0.851	0.011	22.9	35.0	46.2	56.5	1088.3	< 0.001
<i>Forelius pruinosus</i> (Roger, 1863)	0.415	0.797	0.025	6.1	6.5	6.6	6.8	8.3	< 0.001
<i>Linepithema dispertitum</i> (Forel, 1885)	0.350	0.84	0.022	21.5	35.8	44.2	69.8	2858.8	< 0.001
<i>Tapinoma litorale</i> Wheeler, 1905	0.525	0.849	0.016	42.8	61.1	70.6	99.9	479.6	< 0.001
<i>Tapinoma ramulorum</i> Emery, 1896	0.494	0.839	0.012	16.6	18.7	20.3	22.5	40.1	< 0.001
Dorylinae									
<i>Cheliomyrmex morosus</i> (Smith, 1859)	0.365	0.778	0.031	12.1	14.4	16.0	17.6	94.8	< 0.001
<i>Ectiton burchellii</i> (Westwood, 1842)	0.445	0.799	0.009	21.7	24.1	25.1	26.2	34.0	< 0.001
<i>Ectiton mexicanum</i> Roger, 1863	0.335	0.829	0.009	24.0	30.0	34.1	37.5	109.7	< 0.001
<i>Ectiton vagans</i> (Olivier, 1792)	0.377	0.816	0.012	50.8	67.1	75.4	91.6	170.1	< 0.001
<i>Labidus coecus</i> (Latreille, 1802)	0.334	0.778	0.022	5.6	5.9	6.1	6.6	16.2	< 0.001
<i>Labidus praedator</i> (Smith, 1858)	0.243	0.816	0.017	15.0	18.0	19.2	21.5	25.3	< 0.001
<i>Neivamyrmex adhepos</i> (Wheeler, 1922)	0.627	0.975	0.000	52.8	142.8	247.0	377.0	510.9	< 0.001
<i>Neivamyrmex cornutus</i> Watkins, 1975	0.499	0.842	0.033	41.7	91.2	144.9	375.4	713.3	< 0.001
<i>Neivamyrmex halidaii</i> (Shuckard, 1840)	0.632	0.785	0.016	41.3	78.3	97.3	133.6	14796.3	< 0.001
<i>Neivamyrmex harrisii</i> (Haldeman, 1852)	0.301	0.871	0.012	53.4	54.2	65.5	73.6	111.0	0.008
<i>Neivamyrmex impudens</i> (Mann, 1922)	0.479	0.691	0.031	1.6	60.8	133.8	285.8	583.7	< 0.001
<i>Neivamyrmex nigrescens</i> (Cresson, 1872)		0.845	0.016	24.9	29.2	31.4	33.6	65.9	< 0.001
<i>Neivamyrmex opacithorax</i> (Emery, 1894)	0.647	0.856	0.012	18.2	20.6	22.8	40.8	82.5	< 0.001
<i>Neivamyrmex pauxillus</i> (Wheeler, 1903)	0.487	0.769	0.000	1.0	20.1	29.7	91.0	171.5	< 0.001
<i>Neivamyrmex pilosus</i> (Smith, 1858)	0.503	0.65	0.028	4.4	4.9	5.2	7.9	25.7	< 0.001
<i>Neivamyrmex sumichrasti</i> (Norton, 1868)	0.318	0.82	0.016	31.5	194.2	296.1	531.4	970.4	< 0.001
<i>Neivamyrmex swainsonii</i> (Shuckard, 1840)	0.370	0.841	0.013	14.1	16.0	17.1	41.0	70.6	< 0.001
<i>Neivamyrmex tristis</i> (Forel, 1901)	0.597	0.818	0.000	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Nomamyrmex esenbeckii</i> (Westwood, 1842)	0.492	0.777	0.017	15.1	18.6	20.1	24.4	47.6	< 0.001
<i>Syscia augustae</i> (Wheeler, 1902)	0.384	0.102	0.063	0.6	5.1	8.4	16.2	8277.7	< 0.001

Subfamily/Species	10P	TSS	FOR	AUCp					P
				Min	Q1	Q2	Q3	Max	
Ectatomminae									
<i>Ectatomma parasiticum</i> Feitosa & Fresneau, 2008	0.334	1	0.000	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Ectatomma ruidum</i> (Roger, 1860)	0.362	0.787	0.004	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Ectatomma tuberculatum</i> (Olivier, 1792)	0.275	0.803	0.024	12.8	14.6	16.0	17.9	26.2	< 0.001
<i>Gnamptogenys hartmani</i> (Wheeler, 1915)	0.293	0.664	0.038	15.4	32.1	42.7	59.7	172.3	< 0.001
<i>Gnamptogenys striatula</i> Mayr, 1884	0.632	0.818	0.022	3.1	3.2	3.3	4.7	69.3	< 0.001
<i>Gnamptogenys sulcata</i> (Smith, 1858)		0.854	0.012	34.7	53.5	63.8	96.4	333.1	< 0.001
<i>Holcoponera bisulca</i> (Kempf & Brown, 1968)	0.425	0.881	0.010	27.6	40.2	46.0	53.7	87.1	< 0.001
<i>Holcoponera strigata</i> (Norton, 1868)	0.418	0.842	0.010	14.9	16.3	16.9	18.7	46.7	< 0.001
<i>Typhlomyrmex prolatus</i> Brown, 1965	0.482	1	0.000	0.0	0.0	0.0	0.0	0.0	< 0.001
Formicinae									
<i>Acropyga exsanguis</i> (Wheeler, 1909)	0.312	0.818	0.013	10.8	13.7	15.1	32.6	86.9	< 0.001
<i>Brachymyrmex depilis</i> Emery, 1893	0.437	0.823	0.029	3.1	3.2	3.2	3.2	13.5	< 0.001
<i>Brachymyrmex minutus</i> Forel, 1893	0.293	0.822	0.007	19.7	23.0	24.8	27.7	44.0	< 0.001
<i>Brachymyrmex musculus</i> Forel, 1899	0.511	0.812	0.028	63.7	87.9	113.7	211.3	362.0	< 0.001
<i>Brachymyrmex obscurior</i> Forel, 1893	0.347	0.779	0.024	17.4	23.1	24.7	27.0	36.4	< 0.001
<i>Camponotus abscisus</i> Roger, 1863	0.477	0.87	0.015	54.1	72.0	88.5	114.1	7515.9	< 0.001
<i>Camponotus albicoxis</i> Forel, 1899	0.439	0.766	0.020	13.5	102.0	162.7	261.4	8942.8	< 0.001
<i>Camponotus atriceps</i> (Smith, 1858)	0.432	0.768	0.016	11.8	12.9	13.3	13.9	20.5	< 0.001
<i>Camponotus brettesi</i> Forel, 1899	0.292	0.691	0.012	19.4	29.6	39.8	58.8	1361.9	< 0.001
<i>Camponotus claviscapus</i> Forel, 1899	0.418	0.874	0.010	30.2	41.9	50.2	76.5	375.4	< 0.001
<i>Camponotus coloratus</i> Forel, 1904	0.429	0.851	0.000	24.8	40.2	60.7	100.2	2489.6	< 0.001
<i>Camponotus coruscus</i> (Smith, 1862)	0.375	0.918	0.007	20.9	34.5	45.5	271.6	519.7	< 0.001
<i>Camponotus curviscapus</i> Emery, 1896	0.432	0.755	0.022	0.7	42.1	65.9	103.7	2520.8	< 0.001
<i>Camponotus excisus</i> Mayr, 1870	0.279	0.738	0.009	3.2	3.4	3.5	55.1	272.8	< 0.001
<i>Camponotus fasciatellus</i> Dalla Torre, 1892	0.536	0.857	0.000	8.6	23.0	35.0	411877.5	823743.9	< 0.001
<i>Camponotus formiciformis</i> Forel, 1885	0.541	0.9	0.000	8.0	12.5	17.0	26.0	53.0	< 0.001
<i>Camponotus linnaei</i> Forel, 1886	0.345	0.798	0.014	43.9	62.1	70.8	82.9	110.0	< 0.001
<i>Camponotus montivagus</i> Forel, 1885	0.447	0.75	0.083	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Camponotus mucronatus</i> Emery, 1890	0.421	0.927	0.005	86.6	199.6	284.8	456.0	797.6	< 0.001
<i>Camponotus nitidior</i> (Santschi, 1921)	0.381	0.754	0.016	34.7	54.8	72.3	121.9	3810.9	< 0.001
<i>Camponotus novogranadensis</i> Mayr, 1870	0.404	0.829	0.016	3.1	3.2	3.2	3.2	43.3	< 0.001
<i>Camponotus piceatus</i> Norton, 1868	0.601	0.8	0.050	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Camponotus planatus</i> Roger, 1863	0.421	0.779	0.029	3.3	3.3	3.4	7.2	20.4	< 0.001
<i>Camponotus pudorosus</i> Emery, 1925	0.450	0.883	0.019	188.2	481.3	967.0	1222.9	2450.4	< 0.001
<i>Camponotus pullatus</i> Mayr, 1866	0.472	0.793	0.040	33.2	76.0	153.0	175.0	351.0	< 0.001
<i>Camponotus rectangularis</i> Emery, 1890	0.423	0.828	0.018	28.6	36.1	43.7	55.0	125.7	< 0.001
<i>Camponotus rubrithorax</i> Forel, 1899	0.263	0.728	0.054	50.7	117.9	163.6	283.2	5863.2	< 0.001
<i>Camponotus senex</i> (Smith, 1858)	0.352	0.783	0.012	25.7	41.8	48.0	65.5	193.2	< 0.001
<i>Camponotus sericeiventris</i> (Guérin-Méneville, 1838)	0.432	0.802	0.032	9.2	9.8	10.1	24.4	33.2	< 0.001
<i>Camponotus striatus</i> (Smith, 1862)	0.399	0.819	0.014	24.0	30.5	35.3	42.7	74.8	< 0.001
<i>Camponotus textor</i> Forel, 1899	0.324	0.651	0.059	9.0	43.2	62.7	104.1	184.6	< 0.001
<i>Camponotus vicinus</i> Mayr, 1870	0.447	0.897	0.020	3.0	3.1	3.1	3.1	44.9	< 0.001
<i>Camponotus zoc</i> Forel, 1879	0.402	0.629	0.065	3.4	12.5	16.8	23.2	95.5	< 0.001
<i>Colobopsis mathildeae</i> (Smith, M.R., 1949)		0.667	0.167	0.0	0.0	0.0	0.0	0.0	< 0.001

Subfamily/Species	10P	TSS	FOR	AUCp					P
				Min	Q1	Q2	Q3	Max	
Formicinae									
<i>Formica pulla</i> Francoeur, 1973	0.514	1							
<i>Lasius sitiens</i> Wilson, 1955	0.275	0.867	0.015	9.7	11.1	12.0	53.9	690.6	< 0.001
<i>Myrmelachista amicta</i> Wheeler, 1934	0.384	1							
<i>Myrmelachista mexicana</i> Wheeler, 1934	0.372	0.771	0.013	12.4	16.9	20.9	24.6	59.2	< 0.001
<i>Myrmelachista skwarrae</i> Wheeler, 1934	0.295	0.892	0.013	17.1	143.0	345.8	1150.9	2361.5	< 0.001
<i>Myrmelachista zeledoni</i> Emery, 1896	0.304	0.846	0.015	44.2	71.2	94.0	230.4	23131.6	< 0.001
<i>Nylanderia guatemalensis</i> (Forel, 1885)	0.376	0.823	0.016	32.9	49.3	59.3	76.5	135.7	< 0.001
<i>Nylanderia parvula</i> (Mayr, 1870)	0.446	0.858	0.016	3.1	3.1	3.2	72.9	321.6	< 0.001
<i>Nylanderia steinheili</i> (Forel, 1893)	0.496	0.797	0.012	16.3	18.9	26.0	37.2	52.3	< 0.001
Myrmicinae									
<i>Acromyrmex octospinosus</i> (Reich, 1793)	0.457	0.863	0.016	13.3	14.4	15.5	50.4	74.9	< 0.001
<i>Adelomyrmex micans</i> Fernández, 2003	0.434	0.909	0.008	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Adelomyrmex myops</i> (Wheeler, 1910)	0.426	0.881	0.021	33.2	85.2	130.5	229.6	419.4	< 0.001
<i>Adelomyrmex paratristani</i> Longino, 2012	0.523	0.826	0.006	134.1	341.7	470.5	829.7	7648.6	< 0.001
<i>Adelomyrmex robustus</i> Fernández, 2003	0.391	0.686	0.007	350.2	859.2	1338.1	3224.8	6069.3	< 0.001
<i>Adelomyrmex silvestrii</i> (Menozzi, 1931)	0.395	0.841	0.011	35.5	42.8	48.1	94.0	145.3	< 0.001
<i>Adelomyrmex tristani</i> (Menozzi, 1931)	0.307	0.795	0.022	56.6	84.2	95.0	109.7	353.1	< 0.001
<i>Apterostigma pilosum</i> Mayr, 1865	0.479	0.811	0.026	32.2	50.2	86.5	226.7	439.5	< 0.001
<i>Atta cephalotes</i> (Linnaeus, 1758)	0.403	0.836	0.019	14.4	15.9	16.9	18.7	38.0	< 0.001
<i>Atta mexicana</i> (Smith, 1858)	0.338	0.798	0.012	32.8	41.9	45.2	51.7	72.1	< 0.001
<i>Carebara urichi</i> (Wheeler, 1922)	0.311	0.844	0.013	3.1	3.1	3.2	44.4	73.1	< 0.001
<i>Cephalotes minutus</i> (Fabricius, 1804)	0.399	0.821	0.012	19.9	22.9	24.5	27.0	52.6	< 0.001
<i>Cephalotes multispinosus</i> (Norton, 1868)	0.401	0.851	0.015	11.5	15.6	20.0	47.5	139.4	< 0.001
<i>Cephalotes pallens</i> (Klug, 1824)	0.493	0.815	0.024	3.2	3.3	3.5	86.8	481.3	< 0.001
<i>Cephalotes scutulatus</i> (Smith, 1867)	0.368	0.89	0.019	17.4	22.4	26.6	108.0	563.0	< 0.001
<i>Cephalotes setulifer</i> (Emery, 1894)	0.426	0.826	0.022	4.8	96.3	155.1	270.1	28751.3	< 0.001
<i>Cephalotes umbraculatus</i> (Fabricius, 1804)	0.491	0.833	0.023	20.9	28.0	43.8	81.7	292.5	< 0.001
<i>Crematogaster corvina</i> Mayr, 1870	0.477	0.85	0.000	17.0	35.0	35.0	35.0	71.0	0.001
<i>Crematogaster crinosa</i> Mayr, 1862	0.333	0.767	0.020	11.2	11.9	12.7	15.3	34.0	< 0.001
<i>Crematogaster curvispinosa</i> Mayr, 1862	0.456	0.836	0.021	26.8	31.7	36.4	51.5	77.2	< 0.001
<i>Crematogaster erecta</i> Mayr, 1866	0.298	0.818	0.013	31.1	41.1	46.4	53.5	311.7	< 0.001
<i>Crematogaster formosa</i> Mayr, 1870	0.570	0.706	0.032	2.2	89.2	212.8	443.8	922.1	< 0.001
<i>Crematogaster limata</i> Smith, 1858	0.361	0.823	0.016	3.1	3.2	3.2	3.3	33.4	< 0.001
<i>Crematogaster montezumia</i> Smith, 1858	0.386	0.684	0.027	9.9	14.2	18.4	22.7	603.4	< 0.001
<i>Crematogaster nigropilosa</i> Mayr, 1870	0.350	0.872	0.009	8.6	9.4	10.6	19.1	59.6	< 0.001
<i>Crematogaster obscurata</i> Emery, 1895	0.635	0.826	0.010	35.4	63.0	78.5	99.1	5848.2	< 0.001
<i>Crematogaster opaca</i> Mayr, 1870	0.334	0.834	0.023	43.6	73.3	124.1	251.6	6466.1	< 0.001
<i>Crematogaster sumichrasti</i> Mayr, 1870	0.353	0.806	0.007	3.1	3.2	3.2	3.3	79.3	< 0.001
<i>Crematogaster torosa</i> Mayr, 1870	0.379	0.797	0.016	26.6	33.2	35.4	38.3	48.6	< 0.001
<i>Cyphomyrmex costatus</i> Mann, 1922	0.339	0.835	0.013	31.5	44.7	51.8	60.8	165.8	< 0.001
<i>Cyphomyrmex minutus</i> Mayr, 1862	0.232	0.829	0.028	14.8	16.6	17.8	19.2	25.4	< 0.001
<i>Cyphomyrmex rimosus</i> (Spinola, 1851)	0.384	0.838	0.017	14.6	16.2	16.9	17.6	25.5	< 0.001
<i>Cyphomyrmex wheeleri</i> Forel, 1900	0.399	0.796	0.025	3.2	3.4	3.7	5.1	398.1	< 0.001
<i>Eurhopalothrix clypeata</i> Brown & Kempf, 1960	0.421	0.683	0.056	1.0	17.0	36.7	107.0	216.7	< 0.001
<i>Eurhopalothrix hunhau</i> Longino, 2013	0.848	0.818	0.010	0.0	0.0	0.0	0.0	0.0	< 0.001

Subfamily/Species	AUCp								
	10P	TSS	FOR	Min	Q1	Q2	Q3	Max	P
Myrmicinae									
<i>Megalomyrmex drifti</i> Kempf, 1961	0.411	0.778	0.024	3.6	4.1	4.3	37.3	136.0	< 0.001
<i>Megalomyrmex incisus</i> Smith, 1947	0.345	0.805	0.018	44.9	84.1	109.1	143.7	476.5	< 0.001
<i>Monomorium cyaneum</i> Wheeler, 1914	0.475	0.773	0.024	5.3	6.2	6.6	7.1	8.7	< 0.001
<i>Monomorium ebeninum</i> Forel, 1891	0.633	0.8	0.038	33.5	39.4	43.3	47.7	61.8	< 0.001
<i>Monomorium minimum</i> (Buckley, 1867)	0.471	0.862	0.017	3.1	3.1	3.1	7.7	9.2	< 0.001
<i>Mycetomoellerius turrifex</i> (Wheeler, W.M., 1903)	0.254	0.725	0.042	3.2	3.4	3.5	35.3	90.0	< 0.001
<i>Mycetosoritis hartmanni</i> (Wheeler, 1907)	0.318	0.778	0.017	34.8	66.4	84.4	128.6	820.0	< 0.001
<i>Mycocepurus curvispinosus</i> Mackay, 1998	0.336	0.333	0.038	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Mycocepurus smithii</i> (Forel, 1893)	0.248	0.879	0.013	20.8	27.2	31.1	49.7	115.6	< 0.001
<i>Myrmicocrypta dilacerata</i> (Forel, 1885)	0.363	1	0.000	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Nesomyrmex echinatinodis</i> (Forel, 1886)	0.371	0.834	0.017	9.6	12.6	14.2	46.1	232.0	< 0.001
<i>Nesomyrmex wilda</i> (Smith, 1943)	0.541	0.777	0.033	16.8	22.7	28.0	35.5	568.4	< 0.001
<i>Octostruma balzani</i> (Emery, 1894)	0.482	0.856	0.011	11.1	12.3	13.0	31.4	45.1	< 0.001
<i>Octostruma trithrix</i> Longino, 2013	0.460	0.885	0.011	72.9	131.1	221.4	1203.5	7472.2	< 0.001
<i>Paratrachymyrmex intermedius</i> (Forel, 1909)	0.462	0.88	0.012	12.6	24.3	36.0	282.8	553.1	< 0.001
<i>Pheidole absurda</i> Forel, 1886	0.550	0.694	0.033	8.6	11.6	15.0	67.3	708.4	< 0.001
<i>Pheidole agricola</i> Wilson, 2003	0.372	1	0.000	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Pheidole bilimeki</i> Mayr, 1870	0.360	0.719	0.020	11.0	12.2	13.0	22.5	35.3	< 0.001
<i>Pheidole fimbriata</i> Roger, 1863	0.423	0.775	0.017	26.1	33.3	36.9	45.8	102.7	< 0.001
<i>Pheidole flavens</i> Roger, 1863	0.469	0.816	0.007	21.2	24.8	27.5	30.7	38.4	< 0.001
<i>Pheidole gouldi</i> Forel, 1886	0.360	0.853	0.005	24.0	144.2	181.2	237.8	368.4	< 0.001
<i>Pheidole harrisonfordi</i> Wilson, 2003	0.408	0.868	0.006	59.1	68.4	72.9	79.5	98.3	< 0.001
<i>Pheidole insipida</i> Forel, 1899	0.645	0.803	0.030	132.7	471.0	824.9	1886.7	3656.5	< 0.001
<i>Pheidole lamia</i> Wheeler, W.M., 1901	0.416	0.857	0.029	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Pheidole mackayi</i> Wilson, 2003	0.430	1	0.000	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Pheidole nubicola</i> Wilson, 2003	0.317	0.745	0.005	57.7	233.7	351.0	702.9	1406.9	< 0.001
<i>Pheidole oaxacana</i> Wilson, 2003	0.334	0.789	0.011	234.2	502.9	587.1	1006.9	1595.1	< 0.001
<i>Pheidole prattorum</i> Wilson, 2003	0.354	0.849	0.011	33.0	43.2	51.5	64.5	9285.7	< 0.001
<i>Pheidole psilogaster</i> Wilson, 2003	0.398	0.954	0.002	24.8	30.7	36.1	1785.4	3545.5	< 0.001
<i>Pheidole punctatissima</i> Mayr, 1870	0.287	0.823	0.009	3.2	3.3	3.5	18.9	3424.5	< 0.001
<i>Pheidole roushae</i> Wilson, 2003	0.777	0.778	0.013	55.6	88.2	108.0	137.6	413.6	< 0.001
<i>Pheidole simonsi</i> Wilson, 2003	0.553	0.755	0.011	17.0	19.9	22.8	37.9	116.5	< 0.001
<i>Pheidole subarmata</i> Mayr, 1884	0.694	0.808	0.010	22.4	25.6	27.3	29.0	37.4	< 0.001
<i>Pheidole susannae</i> Forel, 1886	0.651	0.783	0.014	3.2	3.3	3.4	20.2	75.7	< 0.001
<i>Pheidole tepicana</i> Pergande, 1896	0.539	0.792	0.021	11.8	14.8	16.2	33.4	58.4	< 0.001
<i>Pheidole titanis</i> Wheeler, 1903	0.419	0.823	0.018	4.9	5.6	6.0	6.6	295.4	< 0.001
<i>Pheidole ursus</i> Mayr, 1870	0.456	0.847	0.009	193.7	337.6	512.0	943.5	28789.2	< 0.001
<i>Pheidole xyston</i> Wilson, 2003	0.420	0.897	0.007	155.4	372.6	629.0	1178.5	17540.5	< 0.001
<i>Pogonomyrmex barbatus</i> (Smith, 1858)	0.269	0.883	0.035	8.0	8.3	8.6	9.6	16.0	< 0.001
<i>Procryptocerus scabriusculus</i> Forel, 1899	0.479	0.845	0.002	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Rhopalothrix weberi</i> Brown & Kempf, 1960	0.203	1	0.000	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Rogeria belti</i> Mann, 1922	0.429	0.843	0.022	23.8	33.3	44.6	232.0	4400.4	< 0.001
<i>Rogeria innotabilis</i> Kugler, 1994	0.417	0.84	0.008	25.1	32.3	38.9	64.5	664.4	< 0.001
<i>Sericomyrmex opacus</i> Mayr, 1865	0.338	0.73	0.021	8.3	39.5	43.9	57.0	754.9	< 0.001
<i>Solenopsis azteca</i> Forel, 1893	0.448	0.865	0.006	20.9	25.6	28.5	33.7	540.7	< 0.001

Subfamily/Species	10P	TSS	FOR	AUCp						P
				Min	Q1	Q2	Q3	Max		
Myrmicinae										
<i>Solenopsis geminata</i> (Fabricius, 1804)	0.384	0.772	0.017	3.3	3.3	3.4	7.0	10.0	< 0.001	
<i>Solenopsis globularia</i> (Smith, 1858)	0.386	0.771	0.020	22.0	27.1	30.0	33.0	91.5	< 0.001	
<i>Solenopsis molesta</i> (Say, 1836)	0.436	0.883	0.016	3.1	3.1	3.2	7.3	44.6	< 0.001	
<i>Solenopsis picea</i> Emery, 1896	0.613	0.793	0.008	20.4	22.5	24.3	26.9	40.6	< 0.001	
<i>Solenopsis succinea</i> Emery, 1890	0.492	0.841	0.011	13.8	27.4	43.2	111.8	3261.7	< 0.001	
<i>Solenopsis tenuis</i> Mayr, 1878	0.397	0.837	0.014	15.6	90.9	155.3	333.9	641.3	< 0.001	
<i>Solenopsis xyloni</i> McCook, 1880	0.257	0.874	0.021	5.8	6.2	6.5	7.6	19.6	< 0.001	
<i>Solenopsis zeteki</i> Wheeler, 1942	0.276	0.795	0.020	37.7	51.6	58.3	71.1	122.8	< 0.001	
<i>Stenamma connectum</i> Branstetter, 2013	0.402	0.913	0.004	3496.8	10492.3	17127.6	35182.0	35182.0	< 0.001	
<i>Stenamma excisum</i> Branstetter, 2013	0.379	0.794	0.008	27.7	46.2	57.6	75.8	417.3	< 0.001	
<i>Stenamma felixi</i> Mann, 1922	0.424	0.891	0.007	101.6	142.3	189.4	267.3	823.7	< 0.001	
<i>Stenamma huachucanum</i> Smith, 1957	0.206	0.798	0.025	4.2	4.6	4.9	33.7	120.9	< 0.001	
<i>Stenamma leptospinum</i> Branstetter, 2013	0.518	0.857	0.000	0.0	0.0	0.0	0.0	0.0	< 0.001	
<i>Stenamma lobinodus</i> Branstetter, 2013	0.665	0.919	0.000	0.0	0.0	0.0	0.0	0.0	< 0.001	
<i>Stenamma maximon</i> Branstetter, 2013	0.384	0.808	0.006	240.9	499.7	674.8	1083.6	64493.8	< 0.001	
<i>Stenamma pelophilum</i> Branstetter, 2013	0.339	0.952	0.000	130.0	522.9	1046.9	2029.2	4190.4	< 0.001	
<i>Stenamma stictosomum</i> Branstetter, 2013	0.442	0.933	0.000	21.9	31.0	52.3	99.0	159.0	< 0.001	
<i>Stenamma vexator</i> Branstetter, 2013	0.393	0.914	0.003	0.0	0.0	0.0	0.0	0.0	< 0.001	
<i>Strumigenys boneti</i> Brown, 1959	0.492	0.772	0.020	21.4	56.4	69.9	106.3	110688.3	< 0.001	
<i>Strumigenys brevicornis</i> Mann, 1922	0.243	0.849	0.008	64.4	78.1	84.4	91.4	118.5	< 0.001	
<i>Strumigenys eggersi</i> Emery, 1890	0.483	0.808	0.029	19.9	22.9	24.3	26.4	37.9	< 0.001	
<i>Strumigenys elongata</i> Roger, 1863	0.648	0.81	0.014	18.8	22.2	23.7	27.0	52.3	< 0.001	
<i>Strumigenys louisianae</i> Roger, 1863	0.453	0.809	0.020	7.5	8.0	8.4	10.7	19.4	< 0.001	
<i>Strumigenys ludia</i> Mann, 1922	0.432	0.885	0.014	23.6	37.9	47.6	93.2	521.3	< 0.001	
<i>Strumigenys margaritae</i> Forel, 1893	0.467	0.871	0.015	3.1	3.1	3.2	3.3	121.0	< 0.001	
<i>Strumigenys mcretes</i> Brown, 1959	0.634	0.818	0.012	32.0	51.8	64.0	152.0	6958.0	< 0.001	
<i>Strumigenys pariensis</i> Lattke & Goitia, 1997	0.255	0.733	0.009	0.6	40.7	46.9	67.1	1825.0	< 0.001	
<i>Strumigenys rogata</i> (Bolton, 2000)	0.347	0.892	0.005	60.2	89.6	110.8	157.7	743.2	< 0.001	
<i>Strumigenys schulzi</i> Emery, 1894	0.461	0.882	0.013	51.0	95.7	138.5	218.9	5546.2	< 0.001	
<i>Strumigenys subedentata</i> Mayr, 1887	0.310	0.845	0.013	26.1	32.5	36.3	43.7	126.2	< 0.001	
<i>Temnothorax aztecus</i> (Wheeler, 1931)	0.485	0.904	0.011	626.0	1891.4	3561.2	5843.2	12021.3	< 0.001	
<i>Temnothorax subditivus</i> (Wheeler, 1903)	0.289	0.79	0.013	3.2	3.3	3.4	19.9	34.2	< 0.001	
<i>Tetramorium spinosum</i> (Pergande, 1896)	0.426	0.827	0.023	9.4	11.0	12.3	25.5	118.3	< 0.001	
<i>Trachymyrmex saussurei</i> (Forel, 1885)	0.406	0.711	0.030	2.7	46.4	63.0	138.7	247.7	< 0.001	
<i>Wasmannia auropunctata</i> (Roger, 1863)	0.419	0.766	0.017	3.2	3.2	3.3	12.1	20.5	< 0.001	
<i>Xenomyrmex panamanus</i> (Wheeler, W.M., 1922)	0.435	1	0.000	0.0	0.0	0.0	0.0	0.0	< 0.001	
Ponerinae										
<i>Anochetus mayri</i> Emery, 1884	0.408	0.842	0.020	29.8	35.7	37.8	42.9	53.1	< 0.001	
<i>Cryptopone gilva</i> (Roger, 1863)	0.311	0.804	0.019	3.1	3.2	3.3	30.5	133.2	< 0.001	
<i>Hypoponera nitidula</i> (Emery, 1890)	0.345	0.872	0.008	3.0	3.1	3.1	24.3	76.1	< 0.001	
<i>Hypoponera opaciceps</i> (Mayr, 1887)	0.442	0.73	0.029	3.2	3.3	3.4	4.8	6.6	< 0.001	
<i>Hypoponera opacior</i> (Forel, 1893)	0.270	0.792	0.022	5.0	5.2	5.3	5.9	9.3	< 0.001	
<i>Hypoponera parva</i> (Forel, 1909)	0.368	0.854	0.012	3.1	3.1	3.2	16.3	111.3	< 0.001	
<i>Leptogenys elongata</i> (Buckley, 1866)	0.363	0.888	0.013	5.6	6.3	6.5	14.6	66.3	< 0.001	
<i>Leptogenys pusilla</i> (Emery, 1890)	0.211	0.88	0.019	34.9	51.5	62.7	92.6	13341.1	< 0.001	

Subfamily/Species	AUCp								
	10P	TSS	FOR	Min	Q1	Q2	Q3	Max	P
Ponerinae									
<i>Leptogenys quiriguana</i> Wheeler, 1923	0.346	0.952	0.000	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Neoponera apicalis</i> (Latreille, 1802)	0.380	0.854	0.019	12.2	14.1	16.4	24.7	65.8	< 0.001
<i>Neoponera carinulata</i> (Roger, 1861)		0.778	0.026	46.8	71.7	90.0	135.5	1305.2	< 0.001
<i>Neoponera crenata</i> (Roger, 1861)	0.495	0.833	0.019	27.7	36.1	48.8	93.3	297.9	< 0.001
<i>Neoponera verenae</i> Forel, 1922	0.218	0.865	0.015	12.2	14.1	15.8	121.0	290.5	< 0.001
<i>Neoponera villosa</i> (Fabricius, 1804)	0.391	0.82	0.019	15.4	16.7	17.5	18.5	30.6	< 0.001
<i>Odontomachus bauri</i> Emery, 1892	0.383	0.82	0.017	12.6	13.8	14.7	18.1	44.9	< 0.001
<i>Odontomachus biolleyi</i> Forel, 1908	0.321	0.889	0.000	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Odontomachus chelifer</i> (Latreille, 1802)	0.423	0.801	0.021	26.8	31.4	34.4	38.5	85.5	< 0.001
<i>Odontomachus laticeps</i> Roger, 1861	0.532	0.835	0.016	16.1	18.9	20.6	27.8	91.1	< 0.001
<i>Odontomachus ruginodis</i> Wheeler, 1908	0.383	0.812	0.022	15.9	19.0	20.8	23.9	44.5	< 0.001
<i>Odontomachus yucatecus</i> Brown, 1976	0.328	0.903	0.007	35.4	48.7	56.9	197.6	354.7	< 0.001
<i>Pachycondyla harpax</i> (Fabricius, 1804)	0.476	0.825	0.017	8.0	8.3	8.7	14.8	20.4	< 0.001
<i>Pachycondyla impressa</i> (Roger, 1861)	0.410	0.831	0.024	30.0	37.1	39.7	45.5	65.7	< 0.001
<i>Platythyrea punctata</i> (Smith, 1858)	0.322	0.844	0.016	18.0	21.0	23.5	33.2	47.6	< 0.001
<i>Ponera exotica</i> Smith, 1962	0.301	0.847	0.009	3.1	3.1	3.2	28.3	55.8	< 0.001
<i>Ponera pennsylvanica</i> Buckley, 1866	0.357	0.872	0.017	3.0	3.1	3.1	14.1	84.5	< 0.001
<i>Pseudoponera stigma</i> (Fabricius, 1804)	0.396	0.779	0.026	18.3	24.0	29.1	46.4	79.0	< 0.001
<i>Rasopone ferruginea</i> (Smith, 1858)	0.475	0.809	0.024	17.9	24.4	28.7	178.1	30625.7	< 0.001
Proceratiinae									
<i>Discothyrea horni</i> Menozzi, 1927	0.437	0.87	0.009	16.9	25.9	30.5	125.6	1758.3	< 0.001
<i>Proceratium mancum</i> Mann, 1922	0.338	0.809	0.022	40.1	88.9	111.5	164.8	1637.2	< 0.001
<i>Proceratium micrommatum</i> (Roger, 1863)	0.267	0.838	0.016	25.9	45.6	75.1	124.8	233.4	< 0.001
Pseudomyrmecinae									
<i>Pseudomyrmex boopis</i> (Roger, 1863)	0.718	0.819	0.019	16.2	18.6	20.3	23.3	57.5	< 0.001
<i>Pseudomyrmex brunneus</i> (Smith, 1877)	0.495	0.704	0.038	4.4	6.0	21.3	58.8	57161.2	< 0.001
<i>Pseudomyrmex caeciliae</i> (Forel, 1913)	0.404	0.72	0.026	0.5	53.1	78.7	112.1	36604.7	< 0.001
<i>Pseudomyrmex championi</i> (Forel, 1899)	0.440	0.861	0.021	60.7	118.6	161.8	274.2	9200.4	< 0.001
<i>Pseudomyrmex cubaensis</i> (Forel, 1901)	0.461	0.763	0.021	35.4	44.8	52.5	58.9	86.5	< 0.001
<i>Pseudomyrmex ejectus</i> (Smith, 1858)	0.464	0.801	0.032	13.5	15.6	17.5	20.6	28.0	< 0.001
<i>Pseudomyrmex elongatus</i> (Mayr, 1870)	0.464	0.768	0.021	3.2	3.3	3.3	3.4	44.1	< 0.001
<i>Pseudomyrmex ferrugineus</i> (Smith, 1877)	0.176	0.878	0.015	34.9	43.2	55.2	86.1	153.5	< 0.001
<i>Pseudomyrmex filiformis</i> (Fabricius, 1804)	0.393	0.799	0.026	25.8	60.4	96.1	144.5	406740.5	< 0.001
<i>Pseudomyrmex gracilis</i> (Fabricius, 1804)	0.326	0.76	0.039	3.2	3.3	3.3	3.3	13.1	< 0.001
<i>Pseudomyrmex ita</i> (Forel, 1906)	0.243	0.862	0.013	22.9	35.0	45.9	69.9	314.9	< 0.001
<i>Pseudomyrmex major</i> (Forel, 1899)	0.361	0.825	0.023	2.1	139.9	215.0	330.3	596.0	< 0.001
<i>Pseudomyrmex oculatus</i> (Smith, 1855)	0.380	0.738	0.004	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Pseudomyrmex pallidus</i> (Smith, 1855)	0.346	0.747	0.030	6.5	6.8	7.1	14.2	19.4	< 0.001
<i>Pseudomyrmex peperi</i> (Forel, 1913)	0.425	0.824	0.018	11.2	15.0	17.6	50.9	115.7	< 0.001
<i>Pseudomyrmex seminole</i> Ward, 1985	0.396	0.66	0.041	3.4	3.7	4.1	6.9	40617.8	< 0.001
<i>Pseudomyrmex simplex</i> (Smith, 1877)	0.361	0.783	0.024	20.9	24.0	26.5	29.5	42.8	< 0.001
<i>Pseudomyrmex spiculus</i> Ward, 1989	0.426	0.818	0.030	0.0	0.0	0.0	0.0	0.0	< 0.001
<i>Pseudomyrmex subater</i> (Wheeler & Mann, 1914)	0.440	0.715	0.022	8.1	46.7	64.9	93.0	34309.3	< 0.001
<i>Pseudomyrmex tenuissimus</i> (Emery, 1906)	0.424	0.852	0.014	9.3	97.1	130.1	181.6	113976.9	< 0.001

Table S2. List of nonnative ant species reported for the Jamapa River watershed in central Veracruz, Mexico.

Subfamily/Species
Dolichoderinae
<i>Tapinoma melanocephalum</i> (Fabricius, 1793); <i>Tapinoma sessile</i> (Say, 1836)
Formicinae
<i>Paratrechina longicornis</i> (Latreille, 1802)
Myrmicinae
<i>Cardiocondyla emeryi</i> Forel, 1881; <i>Cardiocondyla minutior</i> Forel, 1899; <i>Cardiocondyla obscurior</i> Wheeler, 1929; <i>Cardiocondyla venustula</i> Wheeler, 1908; <i>Monomorium floricola</i> (Jerdon, 1851); <i>Monomorium pharaonis</i> (Linnaeus, 1758); <i>Tetramorium bicarinatum</i> (Nylander, 1846); <i>Tetramorium caldarium</i> (Roger, 1857); <i>Tetramorium lanuginosum</i> Mayr, 1870.
Ponerinae
<i>Hypoponera punctatissima</i> (Roger, 1859)

Table S3. Occupancy of ant species in the Jamapa River watershed based on ecological niche modeling and species distribution models. N = number of pixels in a resolution of 30 arc seconds; Percentage (%) represents the percentage of the watershed where the presence of the species was predicted.

Subfamily/Species	N	%	Subfamily/Species	N	%
Amblyoponinae			<i>Neivamyrmex swainsonii</i> (Shuckard, 1840)	12158	95.364
<i>Fulakora orizabana</i> (Brown, 1960)	3272	25.665	<i>Neivamyrmex tristis</i> (Forel, 1901)	11045	86.634
Dolichoderinae			<i>Nomamyrmex esenbeckii</i> (Westwood, 1842)	7925	62.162
<i>Azteca alfari</i> Emery, 1893	9214	72.272	<i>Syscia augustae</i> (Wheeler, 1902)	2703	21.202
<i>Azteca gnava</i> Forel, 1906	5506	43.188	Ectatomminae		
<i>Azteca nigricans</i> Forel, 1899	23	0.18	<i>Ectatomma parasiticum</i> Feitosa & Fresneau, 2008	8456	66.327
<i>Azteca velox</i> Forel, 1899	10959	85.96	<i>Ectatomma ruidum</i> (Roger, 1860)	5260	41.258
<i>Dolichoderus diversus</i> Emery, 1894	6002	47.078	<i>Ectatomma tuberculatum</i> (Olivier, 1792)	3855	30.238
<i>Dolichoderus lutosus</i> (Smith, 1858)	6433	50.459	<i>Gnamptogenys hartmani</i> (Wheeler, 1915)	471	3.694
<i>Dorymyrmex bicolor</i> Wheeler, 1906	8676	68.052	<i>Gnamptogenys striatula</i> Mayr, 1884	8088	63.44
<i>Dorymyrmex insanus</i> (Buckley, 1866)	10358	81.246	<i>Gnamptogenys sulcata</i> (Smith, 1858)	7968	62.499
<i>Forelius damiani</i> Guerrero & Fernández, 2008	11065	86.791	<i>Holcoponera bisulca</i> (Kempf & Brown, 1968)	5517	43.274
<i>Forelius pruinosus</i> (Roger, 1863)	11170	87.615	<i>Holcoponera strigata</i> (Norton, 1868)	8024	62.938
<i>Linepithema dispertitum</i> (Forel, 1885)	1637	12.84	<i>Typhlomyrmex prolatus</i> Brown, 1965	9381	73.582
<i>Tapinoma litorale</i> Wheeler, 1905	11419	89.568	Formicinae		
<i>Tapinoma ramulorum</i> Emery, 1896	3026	23.735	<i>Acropyga exsanguis</i> (Wheeler, 1909)	11268	88.383
Dorylinae			<i>Brachymyrmex depilis</i> Emery, 1893	7230	56.71
<i>Cheliomyrmex morosus</i> (Smith, 1859)	6178	48.459	<i>Brachymyrmex minutus</i> Forel, 1893	3148	24.692
<i>Eciton burchellii</i> (Westwood, 1842)	10381	81.426	<i>Brachymyrmex musculus</i> Forel, 1899	9161	71.857
<i>Eciton mexicanum</i> Roger, 1863	10748	84.305	<i>Brachymyrmex obscurior</i> Forel, 1893	9354	73.37
<i>Eciton vagans</i> (Olivier, 1792)	11232	88.101	<i>Camponotus abscisus</i> Roger, 1863	5018	39.36
<i>Labidus coecus</i> (Latreille, 1802)	6909	54.192	<i>Camponotus albicoxis</i> Forel, 1899	10552	82.767
<i>Labidus praedator</i> (Smith, 1858)	12580	98.674	<i>Camponotus atriceps</i> (Smith, 1858)	11730	92.007
<i>Neivamyrmex adnepos</i> (Wheeler, 1922)	944	7.405	<i>Camponotus brettesi</i> Forel, 1899	8946	70.17
<i>Neivamyrmex cornutus</i> Watkins, 1975	436	3.42	<i>Camponotus claviscapus</i> Forel, 1899	8044	63.095
<i>Neivamyrmex halidaii</i> (Shuckard, 1840)	7066	55.424	<i>Camponotus coloratus</i> Forel, 1904	9376	73.543
<i>Neivamyrmex harrisii</i> (Haldeman, 1852)	7310	57.338	<i>Camponotus coruscus</i> (Smith, 1862)	8187	64.217
<i>Neivamyrmex impudens</i> (Mann, 1922)	11546	90.564	<i>Camponotus curviscapus</i> Emery, 1896	11226	88.054
<i>Neivamyrmex nigrescens</i> (Cresson, 1872)	2635	20.668	<i>Camponotus excisus</i> Mayr, 1870	3461	27.147
<i>Neivamyrmex opacithorax</i> (Emery, 1894)	4445	34.865	<i>Camponotus fasciatellus</i> Dalla Torre, 1892	2019	15.837
<i>Neivamyrmex pauxillus</i> (Wheeler, 1903)	8183	64.185	<i>Camponotus formiciformis</i> Forel, 1885	7794	61.134
<i>Neivamyrmex pilosus</i> (Smith, 1858)	7823	61.362	<i>Camponotus linnaei</i> Forel, 1886	7883	61.832
<i>Neivamyrmex sumichrasti</i> (Norton, 1868)	11897	93.317	<i>Camponotus montivagus</i> Forel, 1885	7	0.055

Subfamily/Species	N	%	Subfamily/Species	N	%
<i>Camponotus mucronatus</i> Emery, 1890	1427	11.193	<i>Crematogaster montezumia</i> Smith, 1858	1779	13.954
<i>Camponotus nitidior</i> (Santschi, 1921)	6681	52.404	<i>Crematogaster nigropilosa</i> Mayr, 1870	9503	74.539
<i>Camponotus novogranadensis</i> Mayr, 1870	364	2.855	<i>Crematogaster obscurata</i> Emery, 1895	4007	31.43
<i>Camponotus piceatus</i> Norton, 1868	10512	82.454	<i>Crematogaster opaca</i> Mayr, 1870	6390	50.122
<i>Camponotus planatus</i> Roger, 1863	8771	68.798	<i>Crematogaster sumichrasti</i> Mayr, 1870	8973	70.382
<i>Camponotus pudorosus</i> Emery, 1925	11657	91.435	<i>Crematogaster torosa</i> Mayr, 1870	9598	75.284
<i>Camponotus pullatus</i> Mayr, 1866	786	6.165	<i>Cyphomyrmex costatus</i> Mann, 1922	4401	34.52
<i>Camponotus rectangularis</i> Emery, 1890	11818	92.697	<i>Cyphomyrmex minutus</i> Mayr, 1862	4312	33.822
<i>Camponotus rubrithorax</i> Forel, 1899	5588	43.831	<i>Cyphomyrmex rimosus</i> (Spinola, 1851)	1490	11.687
<i>Camponotus senex</i> (Smith, 1858)	12434	97.529	<i>Cyphomyrmex wheeleri</i> Forel, 1900	8549	67.056
<i>Camponotus sericeiventris</i> (Guérin-Méneville, 1838)	9005	70.633	<i>Eurhopalothrix clypeata</i> Brown & Kempf, 1960	9225	72.359
<i>Camponotus striatus</i> (Smith, 1862)	11729	91.999	<i>Eurhopalothrix hunhau</i> Longino, 2013	3219	25.249
<i>Camponotus textor</i> Forel, 1899	10882	85.356	<i>Megalomyrmex drifti</i> Kempf, 1961	10019	78.587
<i>Camponotus vicinus</i> Mayr, 1870	9301	72.955	<i>Megalomyrmex incisus</i> Smith, 1947	4971	38.991
<i>Camponotus zoc</i> Forel, 1879	74	0.58	<i>Monomorium cyaneum</i> Wheeler, 1914	344	2.698
<i>Colobopsis mathildeae</i> (Smith, M.R., 1949)	9921	77.818	<i>Monomorium ebeninum</i> Forel, 1891	9799	76.861
<i>Formica pulla</i> Francoeur, 1973	7660	60.083	<i>Monomorium minimum</i> (Buckley, 1867)	9982	78.296
<i>Lasius sitiens</i> Wilson, 1955	221	1.733	<i>Mycetomoellerius turrifex</i> (Wheeler, W.M., 1903)	67	0.526
<i>Myrmelachista amicta</i> Wheeler, 1934	8024	62.938	<i>Mycetosoritis hartmanni</i> (Wheeler, 1907)	7319	57.408
<i>Myrmelachista mexicana</i> Wheeler, 1934	9503	74.539	<i>Mycocoepurus curvispinosus</i> Mackay, 1998	6487	50.882
<i>Myrmelachista skwarrae</i> Wheeler, 1934	9067	71.119	<i>Mycocoepurus smithii</i> (Forel, 1893)	3817	29.94
<i>Myrmelachista zeledoni</i> Emery, 1896	11825	92.752	<i>Myrmicocrypta dilacerata</i> (Forel, 1885)	1339	10.503
<i>Nylanderia guatemalensis</i> (Forel, 1885)	10857	85.16	<i>Nesomyrmex echinatinodis</i> (Forel, 1886)	10401	81.583
<i>Nylanderia parvula</i> (Mayr, 1870)	5	0.039	<i>Nesomyrmex wilda</i> (Smith, 1943)	6167	48.372
<i>Nylanderia steinheili</i> (Forel, 1893)	9231	72.406	<i>Octostruma balzani</i> (Emery, 1894)	7333	57.518
Myrmicinae			<i>Octostruma trithrix</i> Longino, 2013	6877	53.941
<i>Acromyrmex octospinosus</i> (Reich, 1793)	2434	19.092	<i>Paratrachymyrmex intermedius</i> (Forel, 1909)	11521	90.368
<i>Adelomyrmex micans</i> Fernández, 2003	4586	35.971	<i>Pheidole absurda</i> Forel, 1886	1615	12.668
<i>Adelomyrmex myops</i> (Wheeler, 1910)	6272	49.196	<i>Pheidole agricola</i> Wilson, 2003	9466	74.249
<i>Adelomyrmex paratristani</i> Longino, 2012	2853	22.378	<i>Pheidole bilimeki</i> Mayr, 1870	4559	35.76
<i>Adelomyrmex robustus</i> Fernández, 2003	2455	19.256	<i>Pheidole fimbriata</i> Roger, 1863	10553	82.775
<i>Adelomyrmex silvestrii</i> (Menozzi, 1931)	10648	83.52	<i>Pheidole flavens</i> Roger, 1863	9247	72.531
<i>Adelomyrmex tristani</i> (Menozzi, 1931)	6289	49.329	<i>Pheidole gouldi</i> Forel, 1886	8542	67.001
<i>Apterostigma pilosum</i> Mayr, 1865	10125	79.418	<i>Pheidole harrisonfordi</i> Wilson, 2003	12181	95.545
<i>Atta cephalotes</i> (Linnaeus, 1758)	3720	29.179	<i>Pheidole insipida</i> Forel, 1899	3563	27.947
<i>Atta mexicana</i> (Smith, 1858)	7418	58.185	<i>Pheidole lamia</i> Wheeler, W.M., 1901	22	0.173
<i>Carebara urichi</i> (Wheeler, 1922)	9789	76.782	<i>Pheidole mackayi</i> Wilson, 2003	6260	49.102
<i>Cephalotes minutus</i> (Fabricius, 1804)	11062	86.768	<i>Pheidole nubicola</i> Wilson, 2003	10156	79.661
<i>Cephalotes multispinosus</i> (Norton, 1868)	8321	65.268	<i>Pheidole oaxacana</i> Wilson, 2003	12533	98.306
<i>Cephalotes pallens</i> (Klug, 1824)	9762	76.571	<i>Pheidole pratorum</i> Wilson, 2003	4101	32.167
<i>Cephalotes scutulatus</i> (Smith, 1867)	7633	59.871	<i>Pheidole psilogaster</i> Wilson, 2003	5388	42.262
<i>Cephalotes setulifer</i> (Emery, 1894)	8296	65.072	<i>Pheidole punctatissima</i> Mayr, 1870	836	6.557
<i>Cephalotes umbraculatus</i> (Fabricius, 1804)	1195	9.373	<i>Pheidole roushae</i> Wilson, 2003	11616	91.113
<i>Crematogaster corvina</i> Mayr, 1870	12319	96.627	<i>Pheidole simonsi</i> Wilson, 2003	6166	48.365
<i>Crematogaster crinosa</i> Mayr, 1862	10055	78.869	<i>Pheidole subarmata</i> Mayr, 1884	2187	17.154
<i>Crematogaster curvispinosa</i> Mayr, 1862	9115	71.496	<i>Pheidole susannae</i> Forel, 1886	8742	68.57
<i>Crematogaster erecta</i> Mayr, 1866	11165	87.575	<i>Pheidole tepicana</i> Pergande, 1896	8159	63.997
<i>Crematogaster formosa</i> Mayr, 1870	5482	42.999	<i>Pheidole titanis</i> Wheeler, 1903	10445	81.928
<i>Crematogaster limata</i> Smith, 1858	11353	89.05	<i>Pheidole ursus</i> Mayr, 1870	298	2.337

Subfamily/Species	N	%	Subfamily/Species	N	%
<i>Pheidole xyston</i> Wilson, 2003	2669	20.935	<i>Hypoponera opacior</i> (Forel, 1893)	12307	96.533
<i>Pogonomyrmex barbatus</i> (Smith, 1858)	6797	53.314	<i>Hypoponera parva</i> (Forel, 1909)	6874	53.918
<i>Procryptocerus scabriusculus</i> Forel, 1899	3298	25.869	<i>Leptogenys elongata</i> (Buckley, 1866)	10084	79.096
<i>Rhopalothrix weberi</i> Brown & Kempf, 1960	11328	88.854	<i>Leptogenys pusilla</i> (Emery, 1890)	300	2.353
<i>Rogeria belti</i> Mann, 1922	8722	68.413	<i>Leptogenys quiriguana</i> Wheeler, 1923	6	0.047
<i>Rogeria innotabilis</i> Kugler, 1994	3579	28.073	<i>Neoponera apicalis</i> (Latreille, 1802)	9375	73.535
<i>Sericomyrmex opacus</i> Mayr, 1865	5411	42.443	<i>Neoponera carinulata</i> (Roger, 1861)	10035	78.712
<i>Solenopsis azteca</i> Forel, 1893	94	0.737	<i>Neoponera crenata</i> (Roger, 1861)	1619	12.699
<i>Solenopsis geminata</i> (Fabricius, 1804)	7558	59.283	<i>Neoponera verenae</i> Forel, 1922	8344	65.448
<i>Solenopsis globularia</i> (Smith, 1858)	11591	90.917	<i>Neoponera villosa</i> (Fabricius, 1804)	2982	23.39
<i>Solenopsis molesta</i> (Say, 1836)	6750	52.945	<i>Odontomachus bauri</i> Emery, 1892	554	4.345
<i>Solenopsis picea</i> Emery, 1896	236	1.851	<i>Odontomachus bolleyi</i> Forel, 1908	10176	79.818
<i>Solenopsis succinea</i> Emery, 1890	4006	31.422	<i>Odontomachus chelifer</i> (Latreille, 1802)	3151	24.716
<i>Solenopsis tenuis</i> Mayr, 1878	5994	47.015	<i>Odontomachus laticeps</i> Roger, 1861	4815	37.768
<i>Solenopsis xyloni</i> McCook, 1880	6210	48.71	<i>Odontomachus ruginodis</i> Wheeler, 1908	11929	93.568
<i>Solenopsis zeteki</i> Wheeler, 1942	283	2.22	<i>Odontomachus yucatecus</i> Brown, 1976	5165	40.513
<i>Stenamma connectum</i> Branstetter, 2013	1237	9.703	<i>Pachycondyla harpax</i> (Fabricius, 1804)	5995	47.023
<i>Stenamma excisum</i> Branstetter, 2013	6291	49.345	<i>Pachycondyla impressa</i> (Roger, 1861)	10900	85.497
<i>Stenamma felixi</i> Mann, 1922	12747	99.984	<i>Platythyrea punctata</i> (Smith, 1858)	2221	17.421
<i>Stenamma huachucanum</i> Smith, 1957	4214	33.054	<i>Ponera exotica</i> Smith, 1962	5539	43.447
<i>Stenamma leptospinum</i> Branstetter, 2013	7250	56.867	<i>Ponera pennsylvanica</i> Buckley, 1866	1000	7.844
<i>Stenamma lobinodus</i> Branstetter, 2013	10111	79.308	<i>Pseudoponera stigma</i> (Fabricius, 1804)	9875	77.457
<i>Stenamma maximon</i> Branstetter, 2013	11778	92.384	<i>Rasopone ferruginea</i> (Smith, 1858)	8639	67.762
<i>Stenamma pelophilum</i> Branstetter, 2013	3434	26.935	Proceratiinae		
<i>Stenamma stictosomum</i> Branstetter, 2013	8933	70.068	<i>Discothyrea horni</i> Menozzi, 1927	6494	50.937
<i>Stenamma vexator</i> Branstetter, 2013	4752	37.274	<i>Proceratium mancum</i> Mann, 1922	5111	40.089
<i>Strumigenys bonetti</i> Brown, 1959	2518	19.751	<i>Proceratium micrommatum</i> (Roger, 1863)	6416	50.326
<i>Strumigenys brevicornis</i> Mann, 1922	8667	67.982	Pseudomyrmecinae		
<i>Strumigenys eggersi</i> Emery, 1890	7970	62.515	<i>Pseudomyrmex boopis</i> (Roger, 1863)	12711	99.702
<i>Strumigenys elongata</i> Roger, 1863	3526	27.657	<i>Pseudomyrmex brunneus</i> (Smith, 1877)	2988	23.437
<i>Strumigenys louisianae</i> Roger, 1863	4748	37.242	<i>Pseudomyrmex caeciliae</i> (Forel, 1913)	10899	85.489
<i>Strumigenys ludia</i> Mann, 1922	4794	37.603	<i>Pseudomyrmex championi</i> (Forel, 1899)	8715	68.358
<i>Strumigenys margaritae</i> Forel, 1893	7423	58.224	<i>Pseudomyrmex cubensis</i> (Forel, 1901)	9349	73.331
<i>Strumigenys micoetes</i> Brown, 1959	673	5.279	<i>Pseudomyrmex ejectus</i> (Smith, 1858)	9391	73.661
<i>Strumigenys pariensis</i> Lattke & Goitia, 1997	1875	14.707	<i>Pseudomyrmex elongatus</i> (Mayr, 1870)	10915	85.615
<i>Strumigenys rogata</i> (Bolton, 2000)	4326	33.932	<i>Pseudomyrmex ferrugineus</i> (Smith, 1877)	10119	79.371
<i>Strumigenys schulzi</i> Emery, 1894	7937	62.256	<i>Pseudomyrmex filiformis</i> (Fabricius, 1804)	10403	81.599
<i>Strumigenys subedentata</i> Mayr, 1887	715	5.608	<i>Pseudomyrmex gracilis</i> (Fabricius, 1804)	8667	67.982
<i>Temnothorax aztecus</i> (Wheeler, 1931)	11079	86.901	<i>Pseudomyrmex ita</i> (Forel, 1906)	11026	86.485
<i>Temnothorax subditivus</i> (Wheeler, 1903)	4087	32.057	<i>Pseudomyrmex major</i> (Forel, 1899)	5573	43.713
<i>Tetramorium spinosum</i> (Pergande, 1896)	9459	74.194	<i>Pseudomyrmex oculatus</i> (Smith, 1855)	10994	86.234
<i>Trachymyrmex saussurei</i> (Forel, 1885)	7372	57.824	<i>Pseudomyrmex pallidus</i> (Smith, 1855)	6410	50.278
<i>Wasemannia auropunctata</i> (Roger, 1863)	52	0.408	<i>Pseudomyrmex peperi</i> (Forel, 1913)	11707	91.827
<i>Xenomyrmex panamanus</i> (Wheeler, W.M., 1922)	6382	50.059	<i>Pseudomyrmex seminole</i> Ward, 1985	7827	61.393
Ponerinae			<i>Pseudomyrmex simplex</i> (Smith, 1877)	8589	67.37
<i>Anochetus mayri</i> Emery, 1884	1735	13.609	<i>Pseudomyrmex spiculus</i> Ward, 1989	11202	87.866
<i>Cryptopone gilva</i> (Roger, 1863)	9064	71.096	<i>Pseudomyrmex subater</i> (Wheeler & Mann, 1914)	3835	30.081
<i>Hypoponera nitidula</i> (Emery, 1890)	9497	74.492	<i>Pseudomyrmex tenuissimus</i> (Emery, 1906)	9165	71.888
<i>Hypoponera opaciceps</i> (Mayr, 1887)	3494	27.406	<i>Pseudomyrmex viduus</i> (Smith, 1858)	7735	60.671