

Natural history of the critically endangered salamander Pseudoeurycea robertsi.

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Mexico is one of the most diverse countries that is losing a large amount of forest due to land use change, these data put Mexico in fourth place for global deforestation rate, therefore, Mexico occupies the first place in number of endangered species in the world with 665 endangered species. It is important to study amphibians because they are among the most threatened vertebrates on Earth and their populations are rapidly declining worldwide due primarily to the loss and degradation of their natural habitats. Pseudoeurycea robertsi is a micro-endemic and critically endangered Plethodontid salamander from the Nevado de Toluca Volcano and to date almost nothing is known about its natural history therefore, we survey fourteen sites of the Nevado de Toluca Volcano a mountain that is part of the Trans-Mexican Volcanic Belt, Mexico. We carry out the most exhaustive sampling scheme of this species throughout the Nevado de Toluca Volcano to know the number of individuals and the microhabitat features associated with the presence of *P. robertsi*. Likewise, we carry out a morphometric study and coloration measurements of *P. robertsi* individuals and we determine the potential distribution of *P.* robertsi and the other 3 species of pletodontids present in the NTV using ecological niche modeling and to determine the most important habitat features associated with the presence of salamander species, as well as to know the niche overlap among salamander species. This information will help raise conservation strategies for this micro-endemic and critically endangered salamander.

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23 Running title: Natural history of *Pseudoeurycea robertsi*.

- 25 Abstract
- 26 Mexico is one of the most diverse countries that is losing a large amount of forest due to land use
- 27 change, these data put Mexico in fourth place for global deforestation rate, therefore, Mexico
- 28 occupies the first place in number of endangered species in the world with 665 endangered species.
- 29 It is important to study amphibians because they are among the most threatened vertebrates on
- 30 Earth and their populations are rapidly declining worldwide due primarily to the loss and
- 31 degradation of their natural habitats. *Pseudoeurycea robertsi* is a micro-endemic and critically



endangered Plethodontid salamander from the Nevado de Toluca Volcano and to date almost nothing is known about its natural history therefore, we survey fourteen sites of the Nevado de Toluca Volcano a mountain that is part of the Trans-Mexican Volcanic Belt, Mexico. We carry out the most exhaustive sampling scheme of this species throughout the Nevado de Toluca Volcano to know the number of individuals and the microhabitat features associated with the presence of *P. robertsi*. Likewise, we carry out a morphometric study and coloration measurements of *P. robertsi* individuals and we determine the potential distribution of *P. robertsi* and the other 3 species of pletodontids present in the NTV using ecological niche modeling and to determine the most important habitat features associated with the presence of salamander species, as well as to know the niche overlap among salamander species. This information will help raise conservation strategies for this micro-endemic and critically endangered salamander.

43 Key Words: endangered species; conservation; *Plethodontidae*; *Abies* forest; deforestation;
 44 environmental niche modelling; morphometric geometric; *Cyt b*; Nevado de Toluca Volcano;
 45 Trans-Mexican Volcanic Belt.

Introduction

Mexico is one of the most diverse countries that is losing a large amount of forest due to land use change (Masera et al., 1997). Most of the deforestation (80%) is taking place in the central and southern parts of Mexico (Masera et al., 1997; Herrera-Arroyo et al., 2013). These data put Mexico in fourth place for global deforestation rate (FAO, 2006; Ellis and Porter-Bolland, 2008; Heredia-Bobadilla et al., 2016, 2017) this generated that Mexico occupies the first place in number of endangered species in the world with 665 endangered species. Broken down, that includes 71 bird species, 96 mammals, 98 reptiles, 181 fish and 219 amphibians (ICUN, 2019). The Trans-Mexican Volcanic Belt (TMVB) is one of the biogeographic zones with the highest species richness and endemism across the country, and it is the most important region in terms of endemic amphibian and reptile species and the second most important in terms of species number (Flores-Villela and Canseco-Márquez, 2007). However, it is one of the most urbanized areas of the country and this fact is disrupting the natural connectivity of forest landscapes, 3.4% of the TMVB is highly fragmented by urban settlements and 44.7% by agricultural areas (Sunny et al., 2017)



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creating small patches of habitat and reducing metapopulation dynamics between the populations of amphibians' forest (Lowe et al., 2005; Honnay and Jacquemyn, 2007; Herrera-Arroyo et al., 2013).

It is important to study amphibians because they are among the most threatened vertebrates on Earth (Catenazzi, 2015) and their populations are rapidly declining worldwide due primarily to the loss and degradation of their natural habitats (Pimm et al., 1995; Fisher and Shaffer, 1996; Davidson et al., 2001; Marsh and Trenham, 2001; Stuart et al., 2004; Mendelson et al., 2006; Wake and Vredenburg, 2008; Sodhi et al., 2008; Gamble et al., 2007; Ducatez and Shiney, 2017). Mexico is ranked fifth in worldwide amphibian diversity (Ochoa-Ochoa et al., 2009; Ordoñez-Ifarraguerri et al., 2017), however, it is the second worldwide country in number of threatened amphibian species (Frías-Alvarez et al., 2010), with 43% of its 376-species threatened (Parra-Olea et al., 2014), the amphibians are threatened in part because of their low dispersion capacity and small home ranges (Blaustein et al., 1994; Beebee, 2005; Zeisset and Beebee, 2008; Hillman et al., 2014), also, they are highly sensitive to perturbations in both terrestrial and aquatic environments because of their dual life histories, highly specialized physiological adaptations and specific microhabitat requirements (Vitt et al., 1990; Wake, 1991; Blaustein, 1994; Blaustein et al., 1994a; Stebbins and Cohen, 1995). Salamanders are important as top-down controls of many invertebrate species and can also be a source of high energy prey for other predators (Davic and Welsh, 2004). In addition, salamanders can represent an important proportion of the vertebrate biomass in old growth forests (Davic and Welsh, 2004), and are thus vital to ecosystem function, also they provide an important indirect regulatory role in the processing of detritus-litter by ingestion of detritivore prey (Davic and Welsh, 2004).

Roberts' false brook salamander, *Pseudoeurycea robertsi*, is a micro-endemic and critically endangered plethodontid genera/genus found in one of the TMVB volcanoes near Toluca in the state of Mexico: the 'Nevado de Toluca' volcano (NTV; SEMARNAT, 2010; IUCN SSC Amphibian Specialist Group, 2016; Sunny et al., 2019a). *P. robertsi* being one of the most threatened amphibians in Mexico with a vulnerability score of 18 (Wilson et al., 2013) has only been reported in one small area of 8km², within a narrow elevation distribution between 2,900 and 3,600 meters above sea level (masl; IUCN SSC Amphibian Specialist Group 2016). Other reports of *P. robertsi* populations presence within other locations has been confirmed by other studies (see Billie, 2009). Few studies have explored systematic and population genetics of this specie (Lynch



et al., 1983; Parra-Olea, 2002; Bille, 2009; Van Rooij et al., 2011; González-Fernández et al., 2019; Sunny et al., 2019a), more research is needed on its natural history.

Therefore, the aims of this study were: (1) to carry out the most exhaustive sampling scheme of this species throughout the NVT to know the number of individuals and the microhabitat features associated with the presence of *P. robertsi*, (2) to carry out morphometric and coloration measurements of *P. robertsi* individuals, (3) to determine the potential distribution of *P. robertsi* and the other 3 species of pletodontids present in the NTV using ecological niche modeling and to determine the most important habitat features associated with the presence of salamanders species, as well as to know the niche overlap among salamanders species. This information will help raise conservation strategies for this micro-endemic and critically endangered salamander.

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Materials and methods

- 106 Population sampling and species identification
- 107 A total of 150 individuals from 14 populations (Fig. 1) were sampled from mid-April to mid-
- October via visual encounter surveys (Crump and Scott, 1994) across the entire polygon of the
- 109 NTV. Field permits and the approval of the ethics committee were given by the Universidad
- Autónoma del Estado de México and SEMARNAT (SGPA/DGVS/05701/16). During the survey,
- we walked back and forth across transects and looked for salamanders in tree stumps, inside fallen
- logs and under the bark, stones, fallen leaves, and moss. The sampling time was between 9 a.m.
- and 3 p.m., during which an area of approximately 10ha was covered in each sampling site. In
- order to sample salamanders, we took tail samples (2mm) for their DNA extraction (see Sunny et
- al., 2019a for a detailed description) and, subsequently, individuals were released back to the very
- site they were captured/capture site. The tail clipping sampling method is a low impact technique
- that does not affect the survival or growth of salamanders (Arntzen et al., 1999; Polich et al., 2013).
- 118 The salamander tissues were preserved in 90% ethanol and then frozen at -20°C until processing
- the samples.
- For identification in the field, we used the phenotypic characteristics mentioned by Bille, (2009)
- and later with molecular data to corroborate the correct identification of the species. In order to
- achieve the species molecular identification we extracted DNA from the salamanders' tail tips with
- the Vivantis extraction kit, we specifically followed the manufacturer instructions. We used the



extracted DNA as a template for the mitochondrial *cytochrome b* gene (*cyt b*) amplification with the following primers: MVZ15 and MVZ16 (Moritz et al., 1992). Forward and reverse sequences of all individuals were aligned and edited manually using BIOEDIT 7.1.3 (Hall, 1999). The retrieved sequences of *cyt b* were blasted in GenBank (Morgulis et al., 2008) optimized for highly similar sequences (Megablast) to confirm they matched to *Pseudoeurycea robertsi* or *Pseudoeurycea leprosa*, considering the E value and the percentage of identity.

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Morphometric measurements and geometric morphometric

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All P. robertsi individuals were photographed on a millimeter paper to obtain the following morphometric measurements head length (HL), from the tip of the snout to the neck; head width (HW), across the widest point of the head; left eye diameter (LED), wide between eyes (WBE), median body width (MBW), across the trunk midway between the front and hind limb insertions, posterior femur length (PFL), both snout vent length (SVL), tail length (TLO), tail wide (TW) and total length (TL), from the posterior left limb insertion to the tip of the longest outstretched toe. To obtain these morphometric measurements, the pixels of each image were converted to millimeters using the software FIJI (Schindelin et al., 2012). For the geometric morphometric, the shape of the head was obtained using 11 landmarks throughout the contour of this structure, and for the eyes 3 landmarks were located as support points for the layout of fans that covered the entire head structure. For all the analyzed individuals of P. robertsi we used the software MAKEFAN6 (Sheets, 2003) and TPSDIG2 (Rohlf 2005). The configurations of the landmarks of each organism were adjusted by means of Procrustes using the COORDGEN6F (Sheets, 2003) software, in order to eliminate all information not related to the shape. With the software SEMILAND6 (Sheets, 2002), were aligned using the Procrustes superimposition method and the alignment protocol to eliminate the effect of location, orientation and size effect, and thus evaluate the shape variation (Moreno-Barajas et al., 2013). Using this data, we perform a Principal Components Analysis (PCA) using the PCAGEN6P (Sheets, 2002) software to identify the individuals that provide the most intraspecific variation, in addition, the algorithm allowed to visualize the variations of the configuration by means of vectors and deformation grids of each group, which were built applying the thin plate spline interpolation function (Bokstein, 1991). Finally, with the CVAGEN6 (Sheets, 2005) software, an Analysis of Canonical Variables (CVA)



was carried out with a reduction to 2 PC, to identify the possible segregation between groups, depending on the locality.

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Coloration patterns

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- 160 The photographic records were transformed into 8 bits' images and RGB color space. We calculate
- the color frequency of the dorsal stripe of each individual using the 3D Color Inspector and Color
- Histogram plugins (Barthel, 2007) of the ImageJ 1.5 software, and finally we scaled the color
- 163 channels and we made a PCA with the color data.

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165 Microhabitat variables and environmental niche modelling

- When we found a felled log, we captured the geographic coordinates, type of microhabitat, log
- species, if the log was naturally felled, its length and width and whether the log had salamanders,
- we calculated the fallen logs volume, number of individuals in each log and finally, to compare if
- there were differences between sites and characteristics of the microhabitat, different statistical
- tests were performed using the software Paleontological Statistics "PAST" 3.06 (Hammer, 2015).
- 172 To calculated the percentage of forest cover and the percentage of *Abies* cover we used the
- 173 ARCGIS 10.5 software and satellite images (SPOT 6/7) of very high resolution (1.5 m) for year
- 174 2015, we made a supervised classification considering six land cover types: *Pinus* forest, *Abies*
- forest, grasslands, agriculture, urban areas, and water sources. Likewise, to determine the most
- important habitat features associated with the presence of *P. robertsi* we perform an analysis of
- environmental niche modeling (ENM), we also made the ENM for the other species of
- 178 Plethodontids found in the NTV (P. robertsi, P. leprosa, Aquilourycea cephalica and Isthmura
- 179 *bellii*) in order to know if there is competition for environmental space and to know the overlapping
- index with the other Plethodontids. We considered the entire distribution of each species to make
- the ENM. The presence records were obtained from fieldwork and iNaturalist. We selected only
- the records from the last 10 years for the analysis. We defined a polygon (background) for each
- species which represents the accessibility area (Suárez-Atilano et al., 2017; Sunny et al., 2017;
- González-Fernández et al., 2018; Sunny et al. 2019b), considering the biogeographic regions of
- Mexico with geographical records or records near their borders (Sunny et al., 2017; González-



186 Fernández et al., 2018). We obtained bioclimatic variables from WorldClim (Hijmans et al., 2005); topographic and land cover variables were obtained from the National Institute of Statistics and 187 188 Geography. We reclassified the land use map (series V, 1:250000, generated during the period 2011 to 2013; INEGI, 2013) in different exclusive classes that were converted to raster and 189 190 transformed from categorical to continuous using a resample method that averages the value of the surrounding pixels to assign a new value to each pixel (González-Fernández et al., 2018; Sunny et 191 192 al., 2019b). All layers were processed in a raster format, with 1 km resolution, using ARCGIS 10.5 193 and the packages raster (Hijmans, 2016) and rgdal (Bivand et al., 2017) for R software (version 194 3.4.0; R. Development Core Team, 2017). After a bibliographic review and Pearson correlation analysis to discard highly correlated variables ($R^2 > 0.7$, Dormann et al., 2013) we selected the 195 196 following variables: natural grasslands percentage, induced grasslands percentage, percentage of 197 arid vegetation, *Pinus* forest percentage, *Quercus* forest percentage, *Abies* forest percentage, 198 cultivated forest, distance to water sources, agriculture percentage, cloud forest mountain 199 percentage, aquatic vegetation, annual mean temperature, maximum temperature of the warmest month, minimum temperature of the coldest month, precipitation of the wettest month, 200 201 precipitation of the driest month, elevation, aspect, slope (obtained from the elevation layer) and 202 distance to urban settlements, for a better prediction of land use change. We used the maximum 203 entropy modelling software (MAXENT; Phillips et al., 2006). We ran the model for each species in MAXENT with 10 replicates (Anderson et al., 2003; Chefaoui et al., 2005; Suárez-Atilano et 204 205 al., 2017). We used linear and quadratic features because we had less than 80 records in most of 206 the species (Merow et al., 2013). All analyses were performed using the logistic output for an 207 easier interpretation and a convergence threshold of 1 X× 10-5 with 500 iterations (Pearson et al., 2007; Suarez-Atilano 2015; González-Fernández et al., 2018). We generated a continuous map of 208 209 the potential distribution for each species because binary outputs can obscure important biological 210 detail (Liu et al., 2013). To evaluate model performance, we applied the partial ROC analyses 211 (Peterson et al., 2008; Osorio-Olvera et al., 2016). We used the software Niche ToolBox 0.4.1.5 for partial-ROC calculations in R (Osorio-Olvera et al., 2016). 212 213 We compute niche overlap from predictions of species distributions with the "D" similarity 214 statistic of Warren et al., (2008) in the software SDMTools (VanDerWal et al., 2014) for R and we plot the niche overlap from predictions of species distributions in Niche ToolBox. 215



217 Results

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219 Population sampling and species identification

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221 It was quite complicated to differentiate between P. robertsi and P. leprosa in some cases, since 222 the coloration pattern can be confused as Bille, (2009) reported, so the analysis of the species 223 determination using the cyt b confirmed that in all populations 89% of all salamanders found were 224 P. robertsi and 11% were P. leprosa with an E value of 0.0 and an identity percentage of 98% 225 considering a sequence of 617 bp. Some sequences were deposited in the GenBank database with 226 the access numbers: MK357639-MK357709. Finally, we recorded 234 P. robertsi individuals, the largest number of individuals reported for this species. The altitudinal range of our sampling 227 scheme goes from 2800-3460 MASL. At the moment, there is no information on the timing of 228 229 peak of P. robertsi abundance, the months we sampled comprise the beginning of the rainy and 230 dry seasons. In April (beginning of the rainy season) and in October (beginning of the dry season) 231 we did not record any individual. The maximum number of individuals were found at the end of 232 July and the beginning of August, and by the middle of September it was possible to find from 40 233 to 67% less salamanders. The number of individuals within each site in the total sampling scheme 234 varied from 0 to 53, Amanalco A had=53 individuals, Rancho Viejo=29, Meson Viejo=28, 235 Amanalco B=25, Las Lagrimas=24, Carretera=20, Palo Seco=18, Amanalco C=16, Santa 236 Cruz=11, Agua Bendita= 5, Raices= 4, San Juan de las Huertas=1, El Contadero=0 and Huacal 237 Viejo-Agua Bendita=0 (Fig. 2). The 99% of the salamanders were found under the bark of trees, the rest was found under the leaf litter. In Agua Bendita we found together P. robertsi, P. leprosa 238 239 and A. cephalica in two fallen logs (Fig. 2), indicating that these salamanders can share the same 240 environmental niche. We found two individuals of I. bellii in Meson Viejo and one in Rancho 241 Viejo (Fig. 2). We found in August in Palo Seco a *P. robertsi* salamander attending an egg mass 242 of 32 white eggs with transparent parts, the eggs are slightly oval in shape and the clutch was found under the bark of an Abies log. 243

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Morphometric measurements and geometric morphometric

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- 247 The average of the measurements of the juvenile salamanders were: HL=5.475, HW=4.150,
- 248 LED=1.075, WBE=1.800, MBW=4.025, PFL=1.750, SVL=18.543, TLO=14.758, TW=1.975,
- 249 TL=33.300. The average of the measurements of the adult salamanders were: HL=10.795,
- 250 HW=7.288, LED=2.218, WBE=2.712, MBW=7.243, PFL=3.936, SVL=45.414, TLO=43.736,
- 251 TW=3.814, TL=89.150. For the geometric morphometric analysis of the 234 individuals, only 139
- individuals could be analyzed, of which 21 landmarks were placed on the head (Fig. 3A), of which
- 253 six were for the eye contour (Fig. 3A). The deformation grids of the landmarks found a greater
- 254 deformation in the central part of the head (Fig. 3B). The dispersion of the first two main
- components in the PCA analysis showed a single group with a low-medium explained variation
- 256 (45%) between (Fig. 3C). The CVA showed that all individuals of *P. robertsi* from all study areas
- are grouped into a single group (CV1 Eigenvalue= 0.658 and CV2 Eigenvalue=0.491, Axis1:
- 258 λ =0.1604, χ ²=232.431, df=110, p=9.01349 e-011; Axis 2: λ =0.2660, χ ²=168.182, df=90,
- 259 p=1.11354e-006).

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261 Coloration patterns

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- 263 The color of the dorsal stripes ranges from red to orange (R84, G35, B38 to R187 G124 B77; Fig.
- 264 4A), orange to yellow (R174 G135 B63 R191 G178 B118; Fig. 4B) and brown to black (Brown=
- 265 R91 G86 B82 R168 G152 B131 and Black= R28 G28 B28 R68 G68 B68; Fig. 4C). We found
- seven patterns of dorsal stripe coloration, two (Fig. 4a and Fig. 4b) that reported Bille (2009) and
- 5 new patterns (Fig. 4c to 4g), however we could not find the third pattern reported by Bille (2009).
- The patterns that we found were the following: 1) well-defined, brick red dorsal stripe (Fig. 4a),
- 269 2) dense dorsal mottling (Fig. 4b), 3) the dorsal red line only present on the tail (Fig. 4c), 4) semi
- well-defined, yellow dorsal stripe (Fig. 4d), 5) the dorsal yellow line only present on the tail (Fig.
- 4e), 6) almost patternless with few scattered spots (Fig. 4f) and 7) all black without spots or dorsal
- 272 line (Fig. 4g). All these patterns can also have different coloration gradients, with different shades
- of yellow, brown or red.

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275 Microhabitat variables and environmental niche modelling



277 A total of 873 fallen logs and stumps were sampled in all the study areas, the fallen logs volume 278 ranged from 1.3 to 19.6 m³. The sites that only presented *Abies* forest were: El Contadero, Carretera 279 and San Juan de las Huertas (Fig. 5A), the sites that presented the most percentage of total forest 280 in 10 ha were Amanalco A, Amanalco B, Meson Viejo and Rancho Viejo. Likewise, Amanalco 281 A, Amanalco B, Meson Viejo and Rancho Viejo also had the highest percentage of *Abies* forest 282 cover (Fig. 5B). The sites that showed the most percentage of trees naturally fallen were Amanalco 283 A and Meson Viejo, while the sites that presented the most felled logs are Huacal Viejo-Agua Bendita, El Contadero, Carretera, Santa Cruz and Palo Seco (Fig.5C), we found significant 284 differences between the length (F= 8.05 gl= 894 and p=2.08 E-15 Fig. 5D and Fig. 6A) and 285 286 diameter of trees per site (F= 3.359 gl= 894 and p=4.69E-05; Fig. 5D and Fig. 6B) and fallen logs 287 volume per site was greater in Las Lagrimas (Fig. 5E). The average length of the logs with 288 salamanders was 4.2 m and without salamanders was 2.7 m, finding significant differences (t=5.4, 289 gl= 789 and p= 0.0001), the average diameter of logs with salamanders was 35.7 cm and without 290 salamanders was 34.2 cm, no significant differences were found (t=1.02, gl= 789 and p= 0.31), of the logs sampled 76% were Abies and 24% Pinus, the species of tree that presented more 291 292 salamanders was Abies with 77% and Pinus with 23%, we found salamanders in the 75% of Abies 293 fallen logs and 25% of *Pinus* fallen logs, finding significant differences (Chi²=39.98, gl= 13 and 294 p= 0.0001), of these fallen logs 30% were naturally fallen and 70% were logs felled, finding 295 significant differences (Chi²=817.73, gl= 13 and p= 0.0001).

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297 The MAXENT model performed better than expected by random, where the AUC showed high 298 average values across iterations (mean AUC ratios from 0.816 to 0.993 and p-values <0.001; 299 Table. 1). Also, Partial-ROC bootstrap tests showed significant ratio values of empirical AUC over 300 null expectations (mean AUC ratios \geq 1.4 and p-values <0.001 in all cases; Table. 1). The most 301 important habitat features associated with the presence of *P. robertsi* according to the MAXENT 302 model were: Abies forest (38.8%) and maximum temperature of the warmest month (33.4%) for P. leprosa were: maximum temperature of the warmest month (54.1%) and Abies forest (17.9%), 303 304 for A. cephalica were: maximum temperature of the warmest month (42.2%) and Abies forest 305 (21.6%), and for *I. bellii* was: minimum temperature of the coldest month (54.4%). Abies forest cover percentage was positively correlated with probability of presence of P. robertsi, P. leprosa 306 307 and A. cephalica. For I. bellii probability of presence decreased with temperature lower than 5°C.



Probability of presence of *A. cephalica, P. leprosa* and *P. robertsi* decreased as maximum temperature of the warmest month increased. It is important to note that the maximum temperature of the warmest month and *Abies* forest cover percentage were the most important variable to explain the potential distribution of three of the four species. Potential distribution maps for each species are in Figs. 7A–7D. *P. robertsi* is the salamander species with the smallest distribution (Fig. 7A), followed by *P. leprosa* (Fig. 6B), *A. cephalica* (Fig. 7C) and the salamander species that has the widest distribution is *I. bellii* (Fig. 7D), however the four-salamander species are distributed mainly in the mountains of the TMBV, *A. cephalica* is also distributed in the Sierra Madre Oriental and *I. bellii*, in the Sierra Madre Oriental, in the Sierra Madre Occidental, the Sierra Madre del Sur, the Cuenca del Balsas and the Costa Pacifica Mexicana. The greatest niche overlap occurs between *P. robertsi and P. leprosa* (Fig. 8A), followed by *P. robertsi and A. cephalica* (Fig. 8B) and finally between *P. robertsi and I. bellii* (Fig. 8C).

Discussion

Our study reports the first comprehensive evaluation of the natural history of *P. robertsi*, one of the most threatened and extremely endemic salamanders in Mexico. This represents the most extensive dataset recorded in terms of number of samples (N=150), morphological measurements, coloration patterns, localities (11), micro habitat features and macro habitat analyses. Historically there are just one basic ecological study on this species and have considered a significant lesser number of individuals, localities and a very basic habitat information (Bille, 2009). Furthermore, our study is the most complete study of the natural history of this species. This information will help to make informed management decisions focused on the conservation of this micro-endemic species.

Population sampling and species identification

We found that *P. robertsi* was present in all well preserved *Abies* forest of the NTV, presenting a much wider distribution than previously thought (Bille, 2009; IUCN SSC Amphibian Specialist Group, 2016), extending its range of distribution to the entire NVT where the *Abies* forest occurs.

338 The analysis of species identification through *cyt b*, showed that it is complicated to differentiate 339 between *P. robertsi* and *P. leprosa* by the pattern of coloration, size and morphometric



measurements since these characteristics are very variable (Bille 2009), however P. robertsi is 89% more abundant than P. leprosa. The elevation range observed for P. robertsi was slightly larger than previously reported 2,800-3,600 m a.s.l. (Bille, 2009; IUCN SSC Amphibian Specialist Group, 2016), within these altitudinal limits, we corroborate that this species is found in all the Abies forests of the NVT since this forest is distributed in an altitudinal gradient of 2,400 to 3,500 m in the TMVB (Manzanilla-Quiñones et al., 2019). Bille, (2009) reported that at lower altitudes it is more likely to find more individuals of P. leprosa, likewise, in this study, few P. leprosa and A. cephalica individuals were found in all the altitudinal ranges, therefore it is more likely that P. robertsi limits the distribution and abundance of P. leprosa by being a stronger competitor and with higher levels of abundance and higher capacity for sexual encounter than P. leprosa.

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Morphometric measurements and geometric morphometric

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353 Our morphometric measurements are slightly smaller or similar to those reported for *P. leprosa* 354 and A. cephalica (Ramírez-Bautista et al., 2009) however it seems that tail is longer in P. robertsi 355 than in P. leprosa or A. cephalica. In contrast, P. robertsi is smaller than I. bellii (Ramírez-Bautista 356 et al., 2009). All individuals of P. robertsi from the eleven sampling sites presented similar 357 morphometric measurements and the geometric morphometric analysis showed that there are no 358 significant differences between the sampling sites, according to the landmarks used in the head. 359 These results are similar to Baken and Adams' (2019) as they found no differences between 360 salamander species using landmarks in the foot shape (, 2019). This means that salamanders present little morphometric variation between individuals and even between species (Ramírez-362 Bautista et al., 2009; Baken and Adams, 2019), suggesting that, although salamanders have limited 363 dispersal ability, small home ranges (Kleeberger and Werner, 1982; Ovaska, 1988; Gergits and 364 Jaeger, 1990; Gibbs, 1998a, b; Marvin, 1998) and are restricted to forest patches (Petranka et al., 1993; Welsh and Droege, 2001), they can disperse enough to keep gene flow. It has been reported 365 366 that male salamanders have the ability to disperse through harsh habitats (Marsh et al., 2005) 367 especially when there are high levels of competition (Liebgold et al., 2011), this may favor a weak 368 population structure and similar phenotypes between the sampling sites in P. robertsi (Sunny et 369 al., 2019a).



371 Coloration patterns

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In this study, we found two coloration patterns that were previously described by Bille (2009), however we did not find the third pattern that he described, in which the salamander presents a few spots on the body. Considering the patterns described by Bille, (2009) and those found in this study, this species has a total of eight dorsal-tail stripe patterns. The great variety in dorsal-tail stripe patterns and colorations suggests that this species has a great phenotypic variability and we presume there may be more patterns with different types of combinations in addition to those already described. Likewise, we found a great variation in the number, size and shape of the spots and gradients of color. Bille, (2009) reported that the coloration patterns may be correlated with the altitudinal range, however, we do not corroborate this hypothesis, we believed that it may be correlated with the color of the log where they were found.

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Environmental variables and environmental niche modelling

The largest number of individuals was found in well-preserved *Abies* forests (Amanalco A. Amanalco B, Meson Viejo and Rancho Viejo), where they also found more natural fallen Abies and *Pinus* logs. The largest number of individuals was found under the bark of *Abies* logs, because in these trees the bark does not break into small pieces, generating an ecosystem that allows P. robertsi to have shelter and food like: mollusks as slugs, insects and arachnids (Bille, 2009). In low logging sites (Amanalco A and Meson Viejo), we found many fallen logs that fell down with storms. These sites hold the maximum salamander individuals and logs were almost complete (between 5 and 8 meters long approximately). In moderate logging sites, without clearings and cut logs abandoned (between 1 and 3 meters long approximately), there were relatively high individuals also. However, we found few salamanders in forests with many clearings (Huacal Viejo-Agua Bendita, El Contadero, Carretera, Santa Cruz, Palo Seco and Raices), were most of the fallen logs had been extracted or were small and thin. Therefore, we emphasize the need to leave and actively restore the availability of dead wood in managed forests in order to preserve these salamander species (Kouki, 2001; Parra-Olea et al., 2005; Monterrubio-Rico and Escalante-Pliego, 2006; Ramírez-Bautista et al., 2009; Velo-Antón et al., 2013; Medina et al., 2018). Likewise, fallen logs found in the grasslands, in the forest edges or in forest clearings were always completely dry and tough while fallen logs found inside of the forest were all completely wet and



following a process of wood decomposition. We always found salamanders in wet logs with intermediate levels of decomposition (field observations). This is consistent with the importance of forest core areas reported in another study (González-Fernández et al., 2019) and is related with the importance of humidity for this species as Plethodontids do not have lungs and rely on cutaneous respiration (Petranka et al., 1993; deMaynadier and Hunter 1998).

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ENM analysis show that the macro habitat preferred by *P. robertsi* is the *Abies* forest, as well as *P. leprosa* and *A. cephalica*. However, *P. robertsi* is the most *Abies* forest specialist and the species with the most restricted distribution whereas *P. leprosa* and *A. cephalica* are distributed in all the mountains of the TMVB which contain the 91.1% of the *Abies* forests of the country (Sunny et al., 2017). *I. bellii* has a much more extensive distribution in a great variety of vegetation covers. Likewise, whereas *I. bellii* probability of presence decreased with temperature lower than 5°C, *A. cephalica*, *P. leprosa* and *P. robertsi* were limited by maximum temperatures, this can be explained because these species are adapted to high elevations and cold environments. Therefore, temperature increase caused by climate change will affect these three species more than those adapted to lower elevations because species at the highest elevations are limited in their ability to shift distribution upslope, thereby increasing the possibility of becoming extinct (Parra-Olea et al., 2005; Velo-Antón et al., 2013), specially *P. robertsi* for having such a limited distribution. The highest niche overlap occurred between *P. robertsi* and *P. leprosa* followed by *P. robertsi* and *A. cephalica* and finally between *P. robertsi* and *I. bellii*.

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- 423 Conclusions and conservation implications
- Our results suggest that *Abies* forests are extremely important for *P. robertsi*, as well as for *P.*
- 425 leprosa and A. cephalica and that Abies forests with naturally fallen logs will hold more
- 426 salamander individuals. If these forests disappear or suffer grate reductions *P. robertsi* populations
- 427 can diminish or even become extinct. Thus, it is essential to protect the *Abies* forests of the NTV.
- 428 According to some authors *Abies* forest are in the 30.69% of the area in the NTV (16,164.18 ha;
- 429 Franco-Maass et al., 2006), or 6,847.8 ha according to Manzanilla-Quiñones et al., (2019).
- 430 Therefore, it is necessary to carry out a more precise study to delimit the area of the *Abies* forest
- 431 and prevent its deterioration by anthropogenic activities, such as pollution by garbage,
- agrochemicals, fires and deforestation, to prevent the extinction of the populations of *P. robertsi*.



133	However, the Mexican government recently changed the protection status of the NTV from
134	National Park to a less restrictive category in terms of land use. This new category allows forest
135	harvesting practices with commercial proposes as well as the construction of ecotourism centers
136	in almost all Abies forest extension (Fig. 9), which was well preserved under the National Park
137	category (Franco-Maass et al., 2006; Mastretta-Yanes et al., 2014; González-Fernández et al.,
138	2018). Thus, this change of category is compromising the habitat of the endemic and critically-
139	endangered P. robertsi, as well as other forest-dwelling animals (Danaus plexippus, Aquiloeurycea
140	cephalica, Pseudoeurycea leprosa, Isthmura bellii, Sceloporus grammicus, S. spinosus, Aegolius
141	acadicus, Oculatus tolucae, Lynx rufus, among others). Finally, we recommend to priorize in
142	conservation programs the areas of Amanalco, Meson Viejo and Rancho Viejo, since these areas
143	had higher levels of genetic diversity and number of individuals of P. robertsi populations (Sunny
144	et al., 2019a) likewise, these areas also have the most well conserved Abies-Pinus forests of the
145	NTV.
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153	Compliance with ethical standards and conflict of interest
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155	The authors declare that they have no conflict of interest. Our study received the field permits and
156	the approval of the ethics committee from Universidad Autónoma del Estado de México and
157	SEMARNAT (SGPA/DGVS/05701/16).
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159	References
160	
161	Anderson RP, Lew D, Peterson AT. 2003. Evaluating predictive models of species' distributions:
162	criteria for selecting optimal models. Ecological Modelling 162 (3):211–232.

- 463 Arntzen JW, Smithson A, Oldham RS. 1999. Marking and tissue sampling effects on body
- 464 condition and survival in the newt *Triturus cristatus*. Journal of Herpetology 33:567–576.
- 465 Arroyo-Rodríguez V, Saldaña-Vázquez RA, Fahrig L, Santos BA. 2017. Does forest
- fragmentation cause an increase in forest temperature? Ecological Research 32:81–88.
- 467 Baken EK, Adams DC. 2019. Macroevolution of arboreality in salamanders. Ecology and
- Evolution.
- Barthel KU. 2007. Color Inspector 3D. http://rsb.info.nih.gov/ij/plugins/color-inspector.html,
- 470 Accessed: 20019.
- 471 Beebee TJC. 2005. Conservation genetics of amphibians. Heredity 95:423–427.
- 472 Bille T. 2009. Field observations on the salamanders (Caudata: Ambystomatidae, Plethodontidae)
- of Nevado de Toluca, Mexico. Raices 1(2):5.
- Bivand R, Keitt T, Rowlingson B. 2017. rgdal: bindings for the geospatial data abstraction library.
- 475 Blaustein AR, Wake DB, Sousa WP. 1994. Amphibian declines: judging stability, persistence, and
- susceptibility of populations to local and global extinctions. Conservation Biology 8(1):
- 477 60–71.
- 478 Blaustein AR. 1994. Amphibian declines: judging, stability, persistence, and susceptibility of
- populations to local and global extinctions. Conservation Biology 8: 60–71.
- 480 Bookstein FL. 1991. Morphometrics Tools for Landmarks Data: Geometry and Biology.
- 481 Cambridge University Press, Cambridge.
- 482 Catenazzi A. 2015. State of the world's amphibians. Annu Rev Environ Resour 40:91–119.
- 483 Chefaoui RM, Hortal J, Lobo JM. 2005. Potential distribution modelling, niche characterization
- and conservation status assessment using GIS tools: a case study of Iberian Copris species.
- 485 Biological Conservation 122 (2): 327–338.
- 486 Crump ML, Scott NJ Jr. 1994. Visual encounter Surveys. In: Heyer WR, Donnelly MA,
- 487 McDiarmid RW, Hayek LC, Foster MC (eds) Measuring and monitoring biological
- 488 diversity: standard methods for amphibians. Smithsonian Institution Press, Washington, pp
- 489 84–92.
- 490 Davic RD, Welsh Jr HH. 2004. On the ecological roles of salamanders. Annual Review of Ecology,
- Evolution, and Systematics 35: 405–434.
- 492 Davidson C, Shaffer HB, Jennings MR. 2001. Declines of the California red-legged frog: climate,
- 493 UV-B, habitat, and pesticides hypotheses. Ecological Applications 11:464–479.



- 494 deMaynadier PG, Hunter ML. 1998. Effects of silvicultural edges on the distribution and
- abundance of amphibians in Maine. Conservation Biology 12:340–352.
- 496 Dormann CF, Elith J, Bacher S, Buchmann C, Carl G, Carré G, Marquéz JRG, Gruber B,
- Lafourcade B, Leitão PJ, Münkemüller T. 2013. Collinearity: a review of methods to deal
- with it and a simulation study evaluating their performance. Ecography 36 (1): 27–46.
- 499 Ducatez S, Shine R. 2017. Drivers of extinction risk in terrestrial vertebrates. Conservation Letters
- 500 10:186–194.
- 501 Ellis EA, Porter-Bolland L. 2008. Is community-based forest management more effective than
- protected areas?: A comparison of land use/land cover change in two neighboring study
- areas of the Central Yucatan Peninsula, Mexico. Forest Ecology and Management
- 504 256:1971–1983.
- Fahrig L. 2017. Ecological responses to habitat fragmentation per se. Annual Review of Ecology,
- Evolution, and Systematics 48:1–23.
- FAO Statistics database. 2006. WebCite at http://faostat.fao.org/ on 10 June 2016.
- 508 Fisher, RN, Shaffer HB. 1996. The decline of amphibians in California's Great Central Valley.
- 509 Conservation Biology 10:1387–1397.
- 510 Flores-Villela O, Canseco-Márquez L. 2007. Riqueza de la herpetofauna. Universidad Nacional
- Autónoma de México, México, DF: 407–420.
- 512 Franco-Maass S, Regil-García HH, González-Esquivel C, Nava-Bernal G. 2006. Cambio de uso
- del suelo y vegetación en el Parque Nacional Nevado de Toluca, México, en el periodo
- 514 1972–2000. Investicación Geografica 61:38–57.
- 515 Franco-Maass, S, Regil García, HH, Ordóñez-Díaz JAB. 2006. Dinámica de perturbación-
- recuperación de las zonas forestales en el Parque Nacional Nevado de Toluca. Madera y
- 517 Bosques 12(1): 17–28.
- 518 Frías-Alvarez P, Zúniga-Vega JJ, Flores-Villela O. 2010. A general assessment of the
- 519 conservation status and decline trends of Mexican amphibians. Biodiversity and
- 520 Conservation 19(13):3699–3742.
- 521 Gamble LR, McGarigal K, Compton BW. 2007. Fidelity and dispersal in the pond-breeding
- amphibian, *Ambystoma opacum*: implications for spatio–temporal population dynamics
- and conservation. Biological Conservation 139:247–257.

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552

Mexico. Genetica 144:689-698.

524 Gergits WF, Jaeger RG. 1990. Field observations of the behavior of the red-backed salamander 525 (*Plethodon cinereus*): courtship and agonistic interactions. Journal of Herpetology 24(1): 526 93–95. 527 Gibbs JP. 1998a. Amphibian movements in response to forest edges, roads, and streambeds in 528 southern New England. Journal of Wildlife Management 1:584–589. 529 Gibbs JP. 1998b. Distribution of woodland amphibians along a forest fragmentation 530 gradient. Landscape Ecology 13(4):263–268. 531 González-Fernández A, Manjarrez J, García-Vázquez U, D'Addario M, Sunny A. 2018. Present 532 and future ecological niche modelling of garter snake species from the Trans-Mexican 533 Volcanic Belt. PeerJ 6, e4618. González-Fernández A, Arroyo-Rodríguez V, Ramírez-Corona F, Manjarrez J, Aguilera-534 Hernández A, Sunny A. 2019. Local and landscape drivers of the number of individuals 535 536 and genetic diversity of a microendemic and critically endangered salamander. Landscape Ecology 1-12. 537 538 Hall TA. 1999. BioEdit: a user-friendly biological sequence alignment editor and analysis 539 program for Windows 95/98/NT. Nucleic Acids Symposium Series (Vol. 41, No. 41, pp. 95–98). 540 [London]: Information Retrieval Ltd., c1979–c2000. 541 Hammer Ø. 2015. PAST: Paleontological Statistics software package for education and data 542 analysis. Palaeontol Electron 4(1). http://folk.uio.no/ohammer/past/. 543 Helzer CJ, Jelinski DE. 1999. The relative importance of patch area and perimeter-area ratio to 544 grassland breeding birds. Ecological Applications 9:1448–1458. 545 Heredia-Bobadilla RL, Monroy-Vilchis O, Zarco-González MM, Martínez-Gómez D, Mendoza-Martínez GD, Sunny A. 2017. Genetic variability and structure of an isolated population 546 547 of Ambystoma altamirani, a mole salamander that lives in the mountains of one of the largest urban areas in the world. Journal of Genetics 96(6): 873. 548 Heredia-Bobadilla, RL, Monroy-Vilchis O, Zarco-González MM, Martínez-Gómez D, 549 Mendoza-Martínez GD Sunny A. 2016. Genetic structure and diversity in an isolated 550

population of an endemic mole salamander (Ambystoma rivulare Taylor, 1940) of central



553	Herrera–Arroyo ML, Sork	VL,	González–Rodríguez A	A, Rocha–Ramírez V	V ,	Vega I	Ξ, Ο)yama	K.
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- 554 2013. Seed-mediated connectivity among fragmented populations of *Quercus castanea*
- (Fagaceae) in a Mexican landscape. American Journal of Botany 100: 1663–1671.
- 556 Hijmans RJ, Cameron SE, Parra JL, Jones PG, Jarvis A. 2005. Very high resolution interpolated
- climate surfaces for global land areas. International Journal of Climatology 25:1965–1978
- 558 Hijmans RJ. 2016. RASTER (accessed 4 June 2017)
- Hillman SS, Drewes RC, Hedrick MS, Hancok TV. 2014. Physiological vagility: correlations with
- dispersal and population genetic structure of amphibians. Physiological and Biochemical
- 561 Zoology 87: 105–112.
- Honnay O, Jacquemyn H. 2007. Susceptibility of common and rare plant species to the genetic
- 563 consequences of habitat fragmentation. Conservation Biology 21: 823 831.
- 564 INEGI, National Institute for Statistics and Geography. 2013. Conjunto nacional de uso de suelo
- y vegetación a escala 1:250,000 Serie V. México: DGG–INEGI.
- 566 IUCN 2019. The IUCN Red List of Threatened Species. Version 2019–1.
- 567 https://www.iucnredlist.org
- 568 IUCN SSC Amphibian Specialist Group. 2016. Pseudoeurycea robertsi. The IUCN Red List of
- 569 Threatened Species 2016:
- e.T59393A53983925. http://dx.doi.org/10.2305/IUCN.UK.2016–
- 3.RLTS.T59393A53983925.en. Downloaded on 03 May 2018.
- 572 Kapos V, Ganade G, Matsui E, Victoria RL. 1993. 213C as an indicator of edge effects in tropical
- rainforest reserves. Journal of Ecology 81:425–432.
- Kapos, V. 1989. Effects of isolation on the water status of forest patches in the Brazilian Amazon.
- Journal of Tropical Ecology 5(2): 173–185.
- 576 Kleeberger SR, Werner JK. 1982. Home range and homing behavior of *Plethodon cinereus* in
- 577 northern Michigan. Copeia 409–415.
- 578 Kouki J, Löfman S, Martikainen P, Rouvinen S, Uotila A. 2001. Forest fragmentation in
- Fennoscandia: linking habitat requirements of wood-associated threatened species to
- landscape and habitat changes. Scandinavian Journal of Forest Research 16:27–37.
- Liebgold EB, Brodie ED, Cabe PR. 2011. Female philopatry and male-biased dispersal in a
- direct-developing salamander, *Plethodon cinereus*. Molecular Ecology 20(2):249–257.

- 583 Liu C, White M, Newell G. 2013. Selecting thresholds for the prediction of species occurrence
- with presence-only data. Journal of Biogeography 40 (4):778–789.
- Lowe AJ, Boshier D, Ward M, Bacles CFE, Navarro C. 2005. Genetic resource impacts of habitat
- loss and degradation; reconciling empirical evidence and predicted theory for Neotropical
- 587 trees. Heredity 95:255 273.
- 588 Lynch JF, Wake DB, Yang SY. 1983. Genic and morphological differentiation in Mexican
- Pseudoeurycea (Caudata: Plethodontidae), with a description of a new species. Copeia
- 590 884–894.
- 591 Manzanilla-Quiñones U, Martínez-Adriano CA, Aguirre-Calderón ÓA. 2019. Historical and
- current spatial modeling of the sacred fir (Abies religiosa [Kunth] Schltdl. and Cham.) in
- the Trans-Mexican Volcanic Belt. Revista Chapingo Serie Ciencias Forestales y del
- 594 Ambiente 25 (2):201–217.
- Marsh DM, Milam GS, Gorham NP, Beckman NG. 2005. Forest roads as partial barriers to
- terrestrial salamander movement. Conservation Biology 19(6):2004–2008.
- 597 Marsh DM, Trenham PC. 2001. Metapopulation dynamics and amphibian conservation.
- 598 Conservation Biology 15:40–49.
- 599 Marvin GA. 1998. Territorial behavior of the plethodontid salamander *Plethodon kentucki*:
- influence of habitat structure and population density. Oecologia 114(1):133–144.
- 601 Masera OR, Ordoñez MJ, Dirzo R. 1997. Carbon emissions from Mexican forests: Current
- situation and long–term scenarios. Climatic Change 35:265–295.
- Mastretta-Yanes A, Cao R, Nicasio-Arzeta S, Quadri P, Escalante-Espinosa T, Arredondo L,
- Piñero D. 2014. ¿Será exitosa la estrategia del cambio de categoría para mantener la
- biodiversidad del Nevado de Toluca? Oikos 12:7–17.
- Medina JP, Sánchez–Jasso JM, Sealy SG, Salgado–Miranda C, Soriano–Vargas E. 2018. Highest
- Elevational Records for Northern Saw–whet Owls (*Aegolius acadicus*). Journal of Raptor
- 608 Research 52(1):94–100.
- Mendelson JR, Lips KR, Gagliardo RW, Brodie ED. 2006. Confronting amphibian declines and
- extinctions. Science 313:48.
- Merow C, Smith MJ, Silander Jr JA. 2013. A practical guide to MaxEnt for modeling species'
- distributions: what it does, and why inputs and settings matter. Ecography 36 (10):1058–
- 613 1069.

- Monterrubio-Rico TC, Escalante-Pliego P. 2006. Richness, distribution and conservation status
- of cavity nesting birds in Mexico. Biological Conservation 128(1):67–78.
- Moreno-Barajas R, Rodríguez-Romero F, Velázquez-Rodríguez A, Aragón-Martínez A. 2013.
- Variación geográfica en *Phrynosoma orbiculare* (Sauria: Phrynosomatidae): análisis de las
- subespecies. Acta Zoológica Mexicana 29(1):129–143.
- Morgulis A, Coulouris G, Raytselis Y, Madden TL, Agarwala R, Schäffer AA. 2008. Database
- indexing for production MegaBLAST searches. Bioinformatics 24(16):1757–1764.
- 621 Moritz C, Schneider CJ, Wake DB. 1992. Evolutionary relationships within the Ensatina
- 622 eschscholtzii complex confirm the ring species interpretation. Systematic Biology
- 623 41(3):273–291.
- Nowakowski AJ, Watling JI, Thompson ME, Brusch GA, Catenazzi A, Whitfield SM, Kurz DJ,
- Suárez–Mayorga A, Aponte–Gutiérrez A, Donnelly MA, Todd BD. 2018. Thermal biology
- mediates responses of amphibians and reptiles to habitat modification. Ecology Letters
- 627 21:345–355.
- Ochoa-Ochoa L, Urbina-Cardona JN, Vázquez LB, Flores-Villela O, Bezaury-Creel J. 2009. The
- effects of governmental protected areas and social initiatives for land protection on the
- conservation of Mexican amphibians. PLoS One 4(9):e6878.
- Ordoñez-Ifarraguerri A, Siliceo-Cantero HH, Suazo-Ortuño I, Alvarado-Díaz J. 2017. Does a
- Frog Change its Diet along a Successional Forest Gradient? The Case of the Shovel–Nosed
- Treefrog (Diaglena spatulata) in a Tropical Dry Forest in Western Mexico. Journal of
- 634 Herpetology 51(3):411–416.
- Osorio-Olvera L, Barve V, Barve N, Soberón J. 2016. Nichetoolbox: from getting biodiversity
- data to evaluating species distribution models in a friendly GUI environment, R package
- 637 version 0.2. 0.0.
- 638 Ovaska K. 1988. Spacing and movements of the salamander *Plethodon vehiculum*. Herpetologica
- 639 377–386.
- Pandey D. 1993. Forest resources assessment 1990: tropical countries. FAO.
- Parra-Olea G, Flores-Villela O, Mendoza-Almeralla C. 2014. Biodiversidad de anfibios en
- México. Revista Mexicana de Biodiversidad 85:460–466.
- Parra–Olea G, Martínez–Meyer E, De León GPP. 2005. Forecasting Climate Change Effects on
- Salamander Distribution in the Highlands of Central Mexico. Biotropica 37(2):202–208.



- Parra–Olea G. 2002. Molecular phylogenetic relationships of neotropical salamanders of the genus
- Pseudoeurycea. Molecular Phylogenetics and Evolution 22(2):234–246.
- Pearson RG. 2007. Species distribution modelling for conservation educators and practitioners.
- Bulletin of the American Museum of Natural History 3:54–89.
- Peterson AT, Papes M, Soberón J. 2008. Rethinking receiver operating characteristic analysis
- applications in ecological niche modelling. Ecological Modelling 213:63–72.
- Petranka JW, Eldridge ME, Haley KE. 1993. Effects of timber harvesting on southern Appalachian
- salamanders. Conservation Biology 7:363–377.
- Pfeifer M, Lefebvre V, Peres CA, Wearn O, Marsh C, Banks-Leite C, Butchart S, Arroyo-
- Rodríguez V, Barlow J, Cerezo A, Cisneros L, D'Cruze N, Faria D, Hadley A, Klingbeil
- B, Kormann U, Lens L, Rangel GM, Morante–Filho JC, Olivier P, Peters S, Pidgeon A,
- Ribeiro D, Scherber C, Schneider–Maunoury L, Struebig M, Urbina–Cardona N, Watling
- JI, Willig M, Wood E, Ewers R. 2017. Creation of forest edges has a global impact on
- forest vertebrates. Nature 551:187–191.
- Phillips SJ, Anderson RP, Schapire RE. 2006. Maximum entropy modeling of species geographic
- distributions. Ecological Modelling 190:231–259
- 661 Pimm SL, Russell GJ, Gittleman JL, Brooks TM. 1995. The future of biodiversity. Science
- 662 269:347–350.
- Polich RL, Searcy CA, Shaffer E. 2013. Effects of tail clipping on survivorship and growth of
- larval salamanders. Journal of Wildlife Management 77:1420–1425.
- R Development Core Team. 2017. R: a language and environment for statistical computing.
- Vienna: R Foundation for Statistical Computing. (accessed 1 May 2017)
- 667 Ramírez-Bautista A, Hernández-Salinas U, García-Vázquez UO, Leyte-Manrique A, Canseco-
- Márquez L. 2009. Herpetofauna del Valle de México: diversidad y conservación.
- Universidad Autónoma del Estado de Hidalgo, 213.
- 670 Rohlf FJ. 2005. tpsDig2 software. State Univ of New York, Stony Brook, NY.
- 671 Schindelin J, Arganda-Carreras I, Frise E et al. 2012. "Fiji: an open-source platform for
- biological-image analysis", Nature methods 9(7):676–682,
- 673 SEMARNAT. 2010. Norma Oficial Mexicana NOM-059-SEMARNAT-2010, Protección
- ambiental-Especies nativas de México de flora y fauna silvestres-Categorías de riesgo y

- especificaciones para su inclusión, exclusión o cambio. Lista de especies en riesgo. Diario
- Oficial de la Federación, 10 diciembre 2010, México.
- 677 Sheets HD. 2002. Semiland6 software. Canisius College, Buffalo, NY.
- 678 Sheets HD. 2003. IMP-Integrated Morphometrics Package. Buffalo: Department of Physics,
- 679 Canisius College.
- 680 Sheets HD. 2005. CVAGen6, Canonical Variates Analysis program for the analysis of shape,
- based on partial warp scores.
- 682 Sheets HD. 2005. CVAGen6, Canonical Variates Analysis program for the analysis of shape,
- based on partial warp scores.
- 684 Sodhi NS, Bickford D, Diesmos AC, Lee TM, Koh LP, Brook BW, Sekercioglu CH, Bradshaw
- 685 CJA. 2008. Measuring the Meltdown: Drivers of Global Amphibian Extinction and
- Decline. PLoS ONE 3: e1636
- 687 Stebbins RC, Cohen NW. 1995. Declining amphibians. A Natural History of Amphibians.
- Princeton University Press, Princeton, NJ, USA, 210–251.
- 689 Stuart SN, Chanson JS, Cox NA, Young BE, Rodrigues ASL, Fischman DL, Waller RW. 2004.
- Status and trends of amphibian declines and extinctions worldwide. Science 306:1783–
- 691 1786.
- 692 Suárez-Atilano M, Rojas-Soto O, Parra JL, Vázquez-Domínguez E. 2017. The role of the
- 693 environment on the genetic divergence between two Boa imperator lineages. Journal of
- 694 Biogeography 44 (9): 2045–2056.
- 695 Suárez-Atilano M, Rojas-Soto O, Parra JL, Vázquez-Domínguez E. 2017. The role of
- 696 environment on the genetic divergence between two *Boa imperator* lineages. Journal of
- 697 Biogeography 44(9):2045–2056
- 698 Suarez–Atilano M. 2015. Filogeografía de *Boa constrictor* (Serpents: Boidae) en México y Centro
- 699 América. PhD Thesis, Instituto de Ecología, UNAM, México. Available:
- 500 bibliotecacentral.unam.mx. Accessed. 25 November 2015.
- 701 Sunny A, Duarte-deJesus L, Aguilera-Hernández A, Ramírez-Corona F, Suárez-Atilano M,
- Percino-Daniel R, Manjarrez J, Monroy-Vilchis O, González-Fernández A. 2019a.
- Genetic diversity and demography of the critically endangered Roberts' false brook
- salamander (*Pseudoeurycea robertsi*) in Central Mexico. Genetica 1–16.



- 705 Sunny A, Gandarilla-Aizpuro FJ, Monroy-Vilchis O, Zarco-Gonzalez MM. 2019b. Potential
- distribution and habitat connectivity of *Crotalus triseriatus* in Central Mexico. Herpetozoa
- 707 32:139–148.
- 708 Sunny A, González-Fernández A, D'Addario M. 2017. Potential distribution of the endemic
- 709 imbricate alligator lizard (Barisia imbricata imbricata) in highlands of central Mexico.
- 710 Amphibia-Reptilia 38(2): 225–231.
- 711 Temple S. 1986. Predicting impacts of habitat fragmentation on forest birds: a comparison of two
- models. In: Verner J, Morrison M, Ralph CJ (eds) Wildlife 2000: modeling habitat
- relationships of terrestrial vertebrates. University of Wisconsin Press, Madison, Wisconsin,
- 714 USA, pp 301–304.
- 715 Tscharntke T, Tylianakis JM, Rand TA, Didham RK, Fahrig L, Batary P, Bengtsson J, Clough Y,
- 716 Crist TO, Dormann CF, Ewers RM, 2012. Landscape moderation of biodiversity patterns
- and processes-eight hypotheses. Biological Reviews 87(3):661–685.
- Van Rooij P, Martel A, Nerz J, Voitel S, Van Immerseel F, Haesebrouck F, Pasmans F. 2011.
- 719 Detection of Batrachochytrium dendrobatidis in Mexican bolitoglossine salamanders
- using an optimal sampling protocol. EcoHealth 8(2):237–243.
- 721 VanDerWal J, Falconi L, Januchowski S, Shoo L, Storlie C, VanDerWal MJ. 2014. Package
- 722 'SDMTools'. R package.
- 723 VanDerWal J, Falconi L, Januchowski S, Shoo L, Storlie C. 2014. SDMTools: Species
- Distribution Modelling Tools: Tools for processing data associated with species
- 725 distribution modelling exercises. R package version 1:1–221.
- 726 Velo-Antón G, Parra JL, Parra-Olea G, Zamudio KR. 2013. Tracking climate change in a
- dispersal-limited species: reduced spatial and genetic connectivity in a montane
- salamander. Molecular Ecology 22:3261–3278.
- 729 Vitt LJ, Caldwell JP, Wilbur HM, Smith DC. 1990. Amphibians as harbingers of decay.
- 730 BioScience 40(6):418–418.
- Wake DB, Vredenburg VT. 2008. Are we in the midst of the sixth mass extinction? A view from
- the world of amphibians. Proceedings of the National Academy of Sciences of the United
- 733 States of America. 105:11466–11473.
- Wake DB. 1991. Declining amphibian populations. Science 253(5022):860–861.

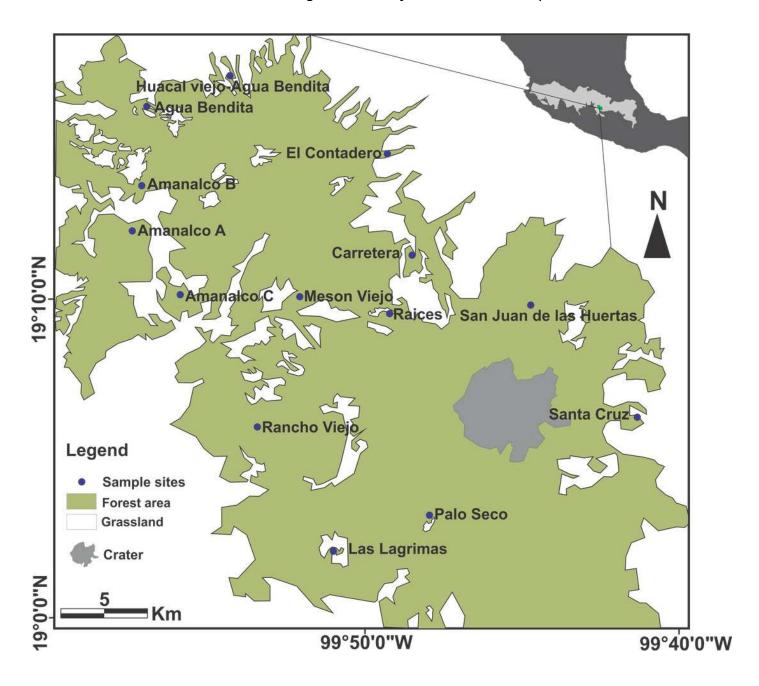


735	Warren DL, Glor RE, Turelli M, Funk D. 2009. Environmental niche equivalency versus
736	conservatism: quantitative approaches to niche evolution. Evolution 62:2868-2883;
737	Erratum: Evolution 65:1215
738	Warren DL, Glor RE, Turelli M. 2008. Environmental niche equivalency versus conservatism:
739	quantitative approaches to niche evolution. Evolution: International Journal of Organic
740	Evolution 6211:2868–2883.
741	Welsh HH, Droege S. 2001. A case for using plethodontid salamanders for monitoring biodiversity
742	and ecosystem integrity of North American forests. Conservation Biology 15(3):558-569.
743	Zeisset I, Beebee TJC. 2008. Amphibian phylogeography: a model for understanding historical
744	aspects of species distributions. Heredity 101:109-119.
745	Zhang Z, Schwartz S, Wagner L, Miller W. 2000. A greedy algorithm for aligning DNA sequences.
746	Journal of Computational Biology 7(1–2):203–14.



Nevado de Toluca Volcano map and sampled sites.

Nevado de Toluca Volcano indicating Pseudoeurycea robertsi sampled sites.

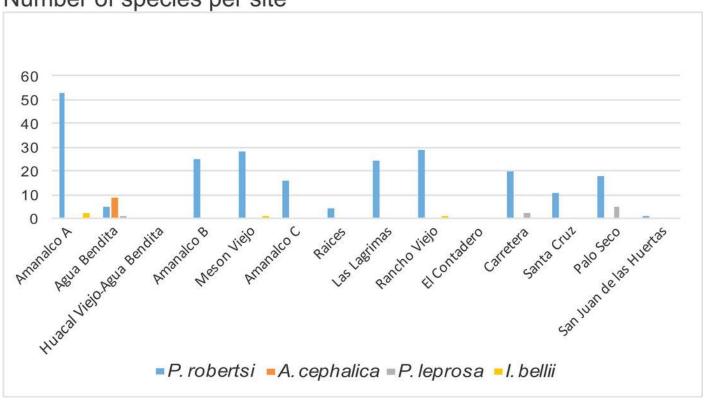




Population sampling

Number of individuals sampled per site.

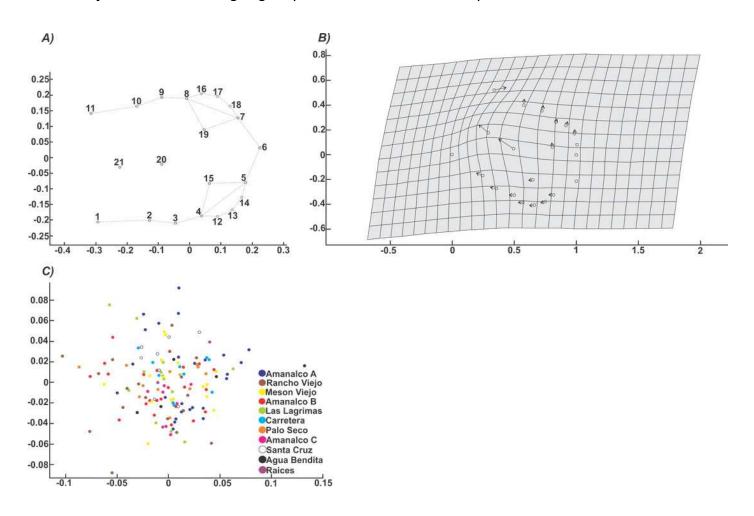
Number of species per site





Morphometric measurements and geometric morphometric

A) Location of landmarks in the head of *Pseudoeurycea robersi*. B) The deformation grid of the landmarks found we found a greater deformation in the central part of the head. C) The PCA analysis showed a single group with a low-medium explained variation.



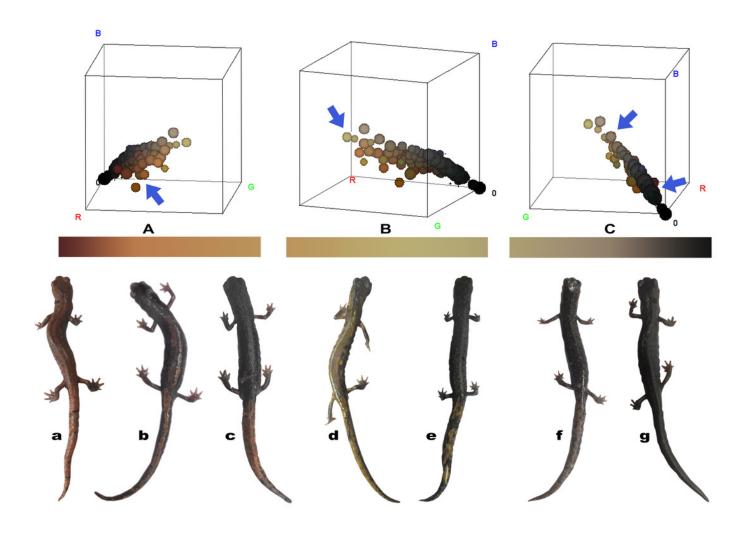


Coloration patterns

Color pattern of the dorsal stripes. A) Dorsal stripes color ranges from red to orange. B)

Dorsal stripes color ranges from orange to yellow. C) Dorsal stripes color ranges from brown to black. The dorsal stripe patterns range from: a) Well-defined, brick red dorsal stripe. b)

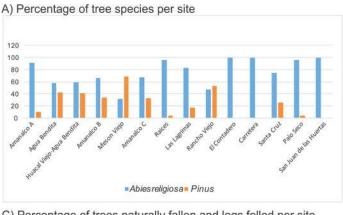
Dense dorsal mottling. c) The dorsal red line only present on the tail. (d) Semi well-defined, yellow dorsal stripe. e) The dorsal yellow line only present on the tail. f) Almost patternless with few scattered spots. g) All black without spots or dorsal line.

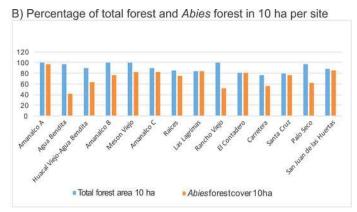


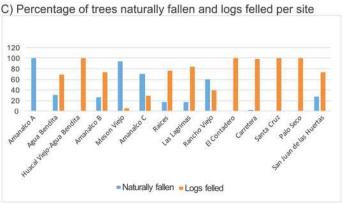


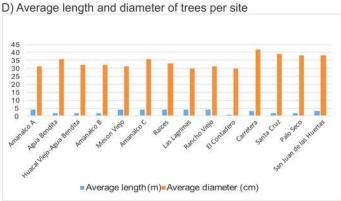
Microhabitat variables

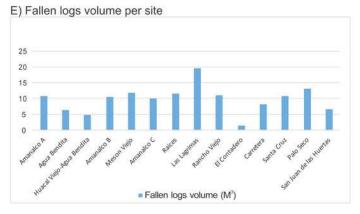
A) Percentage of three species per site. B) Percentage of total forest and *Abies* forest in 10 ha per site. C) Percentage of trees naturally fallen and logs felled per site. D) Average length and diameter of trees per site. E) Falled logs volume per site.







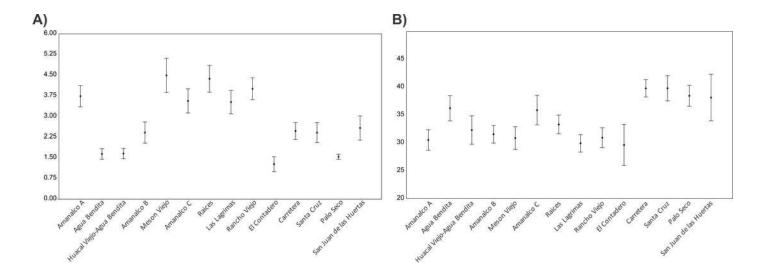






Mean and whisker plots of the ANOVA analysis

Mean and whisker plots of the ANOVA analysis between the A) average length and and B) diameter of trees per site.

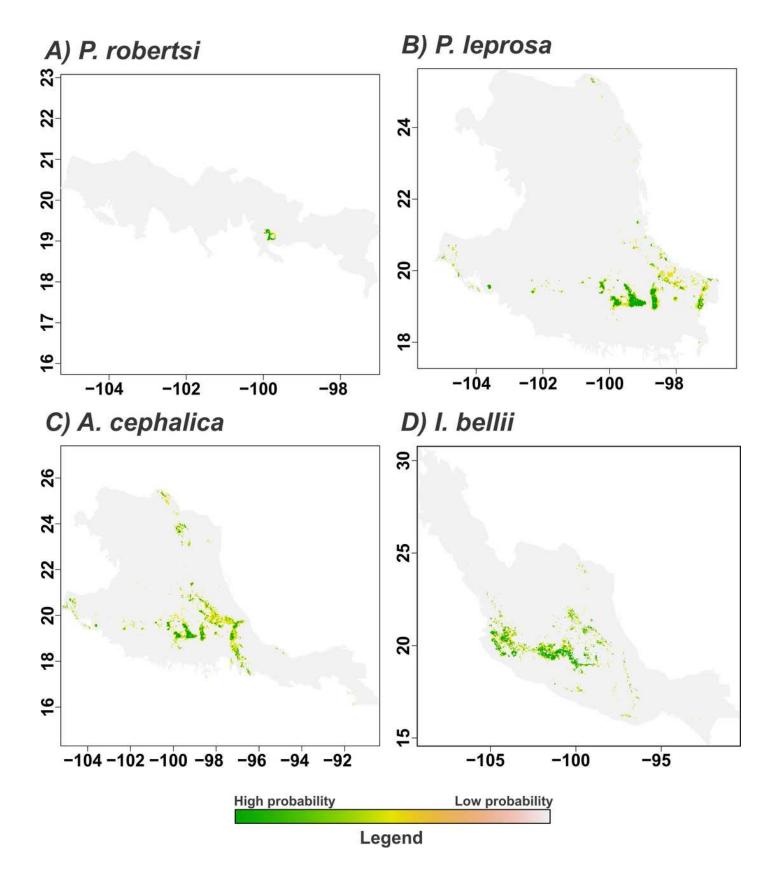




Environmental niche modelling

Potential distribution maps for each salamander species. A). *Pseudoeurycea robersi*. B) *Pseudoeurycea leprosa*. C) *Aquiloeurycea cephalica*. D) and *Isthmura bellii*.





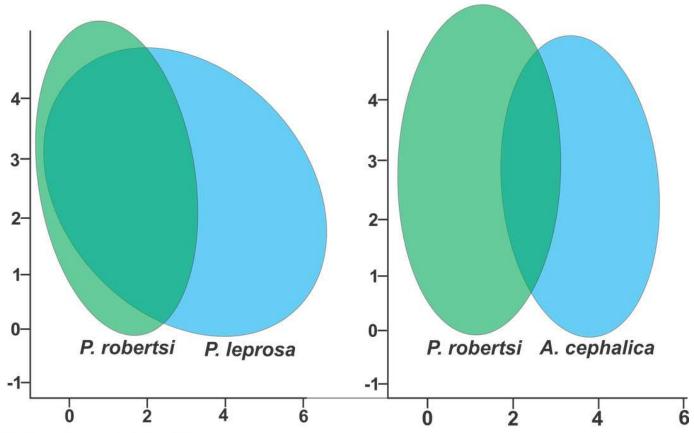


Environmental niche overlap between the salamander species

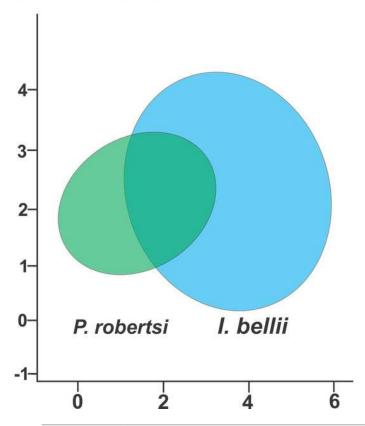
Environmental niche overlap between the four-salamander species. A) Environmental niche overlap between *Pseudoeurycea robersi* and *Pseudoeurycea leprosa*. B) Environmental niche overlap between *Pseudoeurycea robersi* and *Aquiloeurycea cephalica*. C) Environmental niche overlap between *Pseudoeurycea robersi* and *Isthmura bellii*.



B) P. robertsi, A. cephalica



C) P. robertsi, I. bellii





Nevado de Toluca Volcano with the commercial logging allowed areas

Nevado de Toluca Volcano area showing the two types of forest in dark and light green, in red the commercial logging allowed areas and in the blue dots are the sampling sites.

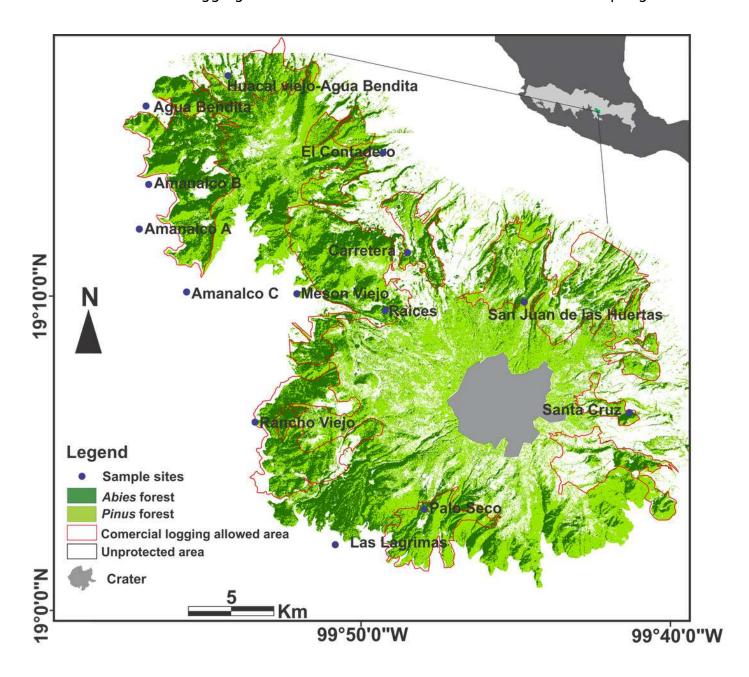




Table 1(on next page)

Potential distribution model performance

Potential distribution AUC and Partial-ROC values.

1

	AUC	Partial-ROC
P. robertsi	0.993	1.970
P. leprosa	0.964	1.846
A. cephalica	0.888	1.430
I. bellii	0.816	1.403