Distribution, richness and conservation of the genus *Salvia* (Lamiaceae) in the State of Michoacán, Mexico

Mayra Flores-Tolentino[‡], Sabina I. Lara-Cabrera[§], José Luis Villaseñor[‡]

- ‡ Universidad Nacional Autónoma de México, Ciudad de México, Mexico
- § Universidad Michoacana de San Nicolás de Hidalgo, Morelia, Mexico

Corresponding author: José Luis Villaseñor (vrios@ib.unam.mx)

Academic editor: Anatoliy Khapugin

Abstract

Little attention has been paid in Mexico to species' geographical distribution, particularly documenting geographic ranges, as a tool to estimate their conservation status. The objective of this study was to review known species distribution and propose potential and conservation status for Salvia species in Michoacán sState using Ecological Niche Models (ENM). We reviewed taxonomic studies for Salvia in Michoacán to compile an initial species checklist, built upon with recently-described species; all the specimens deposited in the National Herbarium were reviewed. The collection data allowed us to build niche models of Salvia species reported for Michoacán. ENM were generated for the species listed using Maxent. In order to minimise collinearity, environmental variables were selected using a Pearson correlation test. Individual models were statistically evaluated and the potential distribution models for each individual species were stacked to obtain the map of richness potential distribution in the State. A total of 66 species of Salvia are listed for Michoacán; however, ENM could only be constructed for 42 of those with ≥ 5 specimens. The environmental variable that most strongly contributed to the models was annual average temperature. The models estimated that Salvia species occupy an area of 23,541 km2 in the State, 72% in the Trans-Mexican Volcanic Belt and a second richest ecoregion is the Sierra Madre del Sur. Although only 3% of the potential distribution area for Salvia in Michoacán is within Protected Areas (PAs), nonetheless, no PA includes rare species. It will therefore be necessary to consider new protection areas or expand existing ones in order to adequately conserve Salvia richness and rarity in the State.

Keywords

conservation, Maxent, ecological niche models, regularisation, species richness

Introduction

Numerous studies document species richness in several regions all over the world; unfortunately, knowledge on the geographic ranges for most of these species is lacking. The few existing reports are often biased by collecting at easily-accessible regions, whereas remote areas are under-sampled. Incomplete sampling has a direct effect on spatial distribution conceptions and negatively influence biogeographic interpretations (Ponder et al. 2001, Whittaker et al. 2005, Boria et al. 2014).

Although much effort has been made to assess species geographic distribution in Mexico, nonetheless, documenting geographical ranges occupied by plants species has been sparse (Villaseñor and Téllez-Valdés 2004). This is unfortunate, given Mexico's high endemism degree (Villaseñor 2016) and the many useful plants (Caballero and Cortés 2001). One such group is *Salvia* (Lamiaceae), with many useful species, but distribution is not well known.

Salvia is the largest genus of the Lamiaceae; worldwide, it is represented by about 1000 species (Walker et al. 2004, Martínez-Gordillo et al. 2017), 306 of which are distributed in Mexico with ca. 77.1% of endemism in the country (Ramamoorthy and Elliot 1998, Martínez-Gordillo et al. 2017). In addition to their species richness and endemism, several species are of economic, medicinal or ornamental importance (Cahill 2003, Moss et al. 2010, Munguía-Lino et al. 2010, Martínez-Gordillo et al. 2013, Akaberi et al. 2016, Martínez-Gordillo et al. 2017). In Mexico, the most emblematic species in traditional medicine include Salvia divinorum Epling & Játiva for its psychoactive activity, Salvia mexicana L. and Salvia tiliifolia Vahl as anti-inflammatory and Salvia hispanica L. as an alimentary supplement (Cahill 2003, Argumedo et al. 2003, Maqueda et al. 2015, González-Chávez et al. 2018). Despite their importance in biodiversity and practical uses, the vast majority of the species' environmental factors that determine their distribution are not known (Martínez-Gordillo et al. 2017).

The genus' taxonomy has been adequately studied for the western Mexican State of Michoacán. Cornejo-Tenorio and Ibarra-Manríquez (2011) reported a total of 64 native species. Later, Lara-Cabrera et al. (2016) in a Lamiaceae-wide list, report only 62 species, through distinct species synonymy conceptions and additions not previously mentioned or recently-described new species. Finally, in a perusal of the Lamiaceae family for Mexico, Martínez-Gordillo et al. (2017) reported 306 species of *Salvia* for the country, 69 distributed in Michoacán. Thus, *Salvia*'s species richness in Michoacán ranges between 62 and 69 species. Several taxonomic studies (Table 1) of infra-generic groups within *Salvia* distributed in the State have also reviewed Michoacán species; *S.* sect. *Sigmoideae* (Espejo-Serna and Ramamoorthy 1993), *S.* sect. *Membranaceae* (González-Gallegos 2014) and *S.* sect. *Scorodoniae* (Olvera-Mendoza et al. 2017). Of equal interest is the taxonomic treatment by González-Gallegos et al. (2016) for the Lamiaceae in the State of Jalisco and species distributed throughout adjacent Michoacán.

The State of Michoacán is amongst the top five most *Salvia*-rich states in Mexico (Cornejo-Tenorio and Ibarra-Manríquez 2011). According to Martínez-Gordillo et al. (2017), six species are endemic to the State (*Salvia cyanantha* Epling, *Salvia gravida* Epling, *Salvia indigocephala* Ramamoorthy, *Salvia madrigalii* Zamudio & Bedolla, *Salvia plurispicata* Epling and *S. synodonta* Epling). *Salvia* species are abundantly distributed in the northeastern region of the State, mainly in temperate and warm regions, at altitudes ranging from 1500 to 3000 m above sea level (Cornejo-Tenorio and Ibarra-Manríquez 2011, Lara-Cabrera et al. 2016). In addition to its great diversity, several *Salvia* species occurring in Michoacán are being investigated for their economic potential in essential oil production, medicinal and ornamental potential (Cornejo-Tenorio and Ibarra-Manríquez 2011, Martínez-Gordillo et al. 2013).

Despite extensive research on the taxonomy of *Salvia* in Michoacán, the rheographic ranges that these species occupy and their relationship with environmental variables has been little explored. Delimiting suitable geographic ranges for species is fundamental, directly underpinning a range of biodiversity and ecosystem function indicators (Jetz et al. 2019), aiding too to traditional morphology-based taxonomy, faced with great challenges in complex taxa (Li et al. 2019). Ecological niche models (ENM) are one of the most commonly used methods for estimating biodiversity patterns, enabling the estimation of species' distributions by association of environmental predictors and presence data (Peterson et al. 2011). In addition, ENM have been widely used in studies dealing with macroecology, conservation, niche evolution, climate change and potential for expansion of invasive species, to mention a few (Hijmans and Graham 2006, Warren et al. 2008, Peterson et al. 2011).

The present research aims to better characterise the distribution of *Salvia* in Michoacán by: 1) updating the known distribution of the genus in the State, 2) using ENM to determine the potential geographic distribution and the environmental variables influencing habitat suitability for these species, 3) comparing the known and potential distribution of the genus to pinpoint areas for further botanic collection and test performance of the ENM and 4) assessing the conservation status both at genus and species level, by analysing habitat loss and coverage of *Salvia* distribution in state-protected natural areas. This approach will identify regions of importance for the conservation of this important genus of the Mexican flora.

Material and methods

Species checklist establishment

We reviewed the most relevant *Salvia* taxonomic papers from Michoacán [Espejo-Serna and Ramamoorthy (1993), Bedolla-García et al. (2011), Cornejo-Tenorio and Ibarra-Manríquez (2011), Iltis et al. (2012), Martínez-Gordillo et al. (2013), González-Gallegos (2014), González-Gallegos et al. (2016), Lara-Cabrera et al. (2016), Martínez-Gordillo et

al. (2017) and Zamudio and Bedolla-García (2018)] and obtained a summarised species list for this study.

Given the taxonomic complexities of the group, with frequent synonymy changes, we only considered the species that were reported in at least two of those studies and that could be corroborated with herbarium specimens. Discrepancies in the *Salvia* list include the number of species, 78 in three papers (Cornejo-Tenorio and Ibarra-Manríquez 2011, Lara-Cabrera et al. 2016, Martínez-Gordillo et al. 2017), although this number is slightly overestimated; for example, *Salvia arbuscula* Fernald is recognised by Cornejo-Tenorio and Ibarra-Manríquez (2011), but considered as a synonym to *S. iodantha* Fernald by Lara-Cabrera et al. (2016). Martínez-Gordillo et al. (2017) do not even include the species for the State. Finally, we updated the list including a recently-described species *S. madrigalii* Zamudio and Bedolla-García (2018).

Occurrence data

Salvia occurrences information for Michoacán was obtained from databases of the National Biodiversity Information System of Mexico (SNIB-REMIB) of the Comisión Nacional para el Conocimiento y Uso de la Biodiversidad (CONABIO), the digital repository of the National Herbarium of Mexico (MEXU-UNIBIO) of the Instituto de Biología, Universidad Nacional Autónoma de México (UNAM) and through the review of taxonomic studies. All specimens deposited in the National Herbarium (MEXU) were reviewed, verifying species identification (see Suppl. material 1).

Cleaning of database

The database was corrected following the recommendations by Chapman (2005) and Castillo et al. (2014) with the following procedure: 1) correct identification of the specimens as verified by specialists, 2) eliminate synonym and duplicates, 3) geo-referencing the localities lacking this information in the specimen label, in Google Earth (https://www.google.com/earth/), using locality name and description and finally 4) eliminate correlated records through pattern analysis.

Pattern analysis was applied to the data for all species using the Ilwis v.3.4 programme (http://52north.org/ilwis). This analysis allowed us to estimate the distance at which the collecting points are not correlated (Cruz-Cárdenas et al. 2014a). The distance obtained from the pattern analysis was used to filter the uncorrelated data through the R *spThin* package (Aiello-Lammens et al. 2015, R Development Core Team 2016), integrated in the Data module Process Occurrence in Wallace v.1.0.6 (Kass et al. 2018). This analysis reduced sampling *bias* by eliminating records per geographic distance, which significantly influences the models performance (Boria et al. 2014).

In addition to the review of the specimens in the herbarium, specimens cited in taxonomic works of the study group were also considered. For the species and specimens with taxonomic circumscription issues, the specimens were individually evaluated to determine whether or not to include them in the analyses. The outliers identified in the pattern

analysis were reviewed to confirm the identity from a genus specialist; the specimens that were not approved by the specialist, were excluded from the database.

Known distribution

The polygon for the state of Michoacán was divided into 118 grid-squares of 15' latitude and 15' longitude; a finer scale grid would have resulted in no species being represented. Thus 15' x 15' grid was a compromise between the distribution, collection effort and the number of grids. Geographic information was analysed in ArcGIS 10.2 (ESRI 2013) to produce a known distribution map and intersecting the grids with the number of species recorded in each grid. Grids were subsequently grouped per species number.

Environmental data

Fifty eight environmental variables were considered, with a resolution of 30 arc seconds, a pixel size of about 1 km 2 (Table 2); 26 of them were climatic (Hijmans et al. 2005), nine topographic, nine edaphic and 14 included remote sensing data (Cruz-Cárdenas et al. 2014b). However, it is known that the use of a large number of variables influences the final results of the models, making it necessary to select only those that are most significant (Peterson and Nakazawa 2008, Varela et al. 2014). To reduce the total number of variables included in models and to minimise multicollinearity amongst the 58 variables, their correlation values (Pearson) were evaluated using NicheToolsBox (Osorio-Olvera et al. 2018); uncorrelated variables were included in the analysis and correlated variables (r > 0.85) were excluded from the analysis.

Ecological niche models

Species distribution result from several factors; amongst the most important ones are environmental variables (A), biotic component (B) and the set of sites that has been accessible to the species (M). Soberón and Peterson (2005) illustrated the interaction of these factors in a diagram called BAM. We used the WWF Ecoregions following Olson et al. (2001) to delimit the accessible area for all species (M in the BAM diagram, Soberón and Peterson 2005), considering only the ecoregions with occurrence records of *Salvia* in Michoacán. Selected ecoregions were then cut, using the State's polygon as a limit.

Evaluating models using spatially independent data improves model configurations and balances their complexity. There are several methods for identifying optimal model configuration (Muscarella et al. 2014, Hijmans et al. 2017). Here, we used the package ENMeval v.0.1.2 in R (Muscarella et al. 2014) to determine the best performance models, run with multiple regularisation (MR) values, from 1 to 4 (in a 1 increment), with four different feature class combinations (L, LQ, H, LQH, where L = Linear, Q = Quadratic, H = Hinge). ENMeval provides multiple evaluations that allow the identification of the optimal model configuration (Muscarella et al. 2014).

Maxent 3.4.1 software (Phillips et al. 2017) was used to run the ENM. Maxent has gained popularity for ENM of many plant and animal species due to its simplicity in configuration. Models were constructed by changing the regularisation values and the feature class, following the results from ENMeval analysis. The default options of clamping and extrapolate were omitted, in order to avoid extrapolation of the extreme values of variables (Elith et al. 2011). Software was set to use 75% of the data to run the analysis and 25% to validate the model (Phillips et al. 2006), only for species with more than five records, the minimum number required in Maxent (Hernandez et al. 2006). For a few records species (5-20), we decided to construct the models using the basic configuration of Maxen (Phillips et al. 2017) and not the feature classes and regularisation multiplier, since these generally result in low Area under the Curve (AUC) values and overprediction. Species with less than five records were considered for generating the map of known richness and are considered amongst the rare species.

Results from Maxent were then processed in ArcGIS 10.2 (ESRI 2013) to obtain binary maps, using the maximum training sensitivity plus specificity as the cutoff threshold (Liu et al. 2016). These binary maps were added with the Algebra map tool to obtain a richness map of *Salvia* species. In addition, the known richness in the State was obtained from collection records in digitised databases, compared with the potential richness obtained through the models. Mexdem (digital elevation model) altitude layer in ArcGIS was used to extract the altitude values of each record of *Salvia* species.

Training/testing partitioning and model evaluation

We partitioned the data into training and testing groups (k-fold cross validation), allowing us to evaluate models' performance (Peterson et al. 2011). We chose to use "block (k-4)", one of the options implemented in ENMeval (Muscarella et al. 2014), which divides the presence data according to its longitude and latitude. This method is one of the most recommended, since it provides better spatial independence of the data (Radosavljevic and Anderson 2014, Fourcade et al. 2018).

The models were evaluated using the AUC, a value that is part of Maxent's results. AUC values range from 0 to 1, where values close to 1 indicate models with perfect discrimination ability and values less than 0.5 indicate that the model is no better than a randomly-generated model (Peterson et al. 2011, Fourcade et al. 2018). Individual models were also evaluated by the binomial test analysis and partial-ROC (receiver operating characteristic), considering a significance value < 0.05. This allowed the evaluation of the models' ability to predict the largest possible number of independent points amongst those that were not used to create the model. The tests were carried out using the R statistical package and NicheToolBox (Osorio-Olvera et al. 2018).

Conservation status

To assess the conservation status of the genus *Salvia*, habitat loss and the area currently within a Protected Area (PA) were analysed along the potential distribution of the genus in

the State of Michoacán. To analyse the habitat loss of the genus, a land use map of Series I and VI of the National Institute of Geography and Informatics (INEGI 1997, INEGI 2017) was used. Maps were then reclassified into two categories: vegetated and non-vegetated (Staude et al. 2019); (1) Specifically, those areas that represent some forest cover within the State polygon were selected (e.g. oak forest, coniferous forest, evergreen forest, deciduous forest, sub-deciduous forest and xerophytic scrub). (2) To estimate the degree of protection of *Salvia*, the map of potential distribution was overlapped with the map of protected natural areas of Mexico (CONANP 2010).

Results

Species and occurrence data

A list of 66 species was obtained for the Mexican State of Michoacán (Suppl. material 2) from bibliographic review of *Salvia*. Fifty two species are reported as endemic to Mexico and three species are restricted to the State (*S. madrigalii*, *S. subobscura* and *S. synodonta*). A complete dataset record of 3,093 was later reduced to 1,925 records after data cleaning. Of these records, 1,836 correspond to the 42 *Salvia* species with pattern analysis and, finally, 404 records were used to build the ENMs.

Patterns of distribution of Salvia in Michoacán

Known distribution

The political territory of Michoacán comprise five ecoregions; *Salvia* has been reported in all (Fig. 1). Of these the Trans-Mexican Volcanic Belt (TMVB) contained the largest number of species (56 species), followed by 21 species in the Sierra Madre del Sur (SMS) and 14 in the Jalisco Dry Forest (JDF).

The Salvia endemic species to the state occupy a very restricted area, as exemplified by S. madrigalii, having being reported at only three sites in the Morelia Municipality, in northeastern Michoacán. A similar case is S. subobscura in the south, known from only two locations in the Chinicuila Municipality. Eight in the restricted species group are here regarded as rare species (< 3 records), known only from one locality (the type locality, Fig. 1), five from western Michoacán, in Chinicuila Municipality (Salvia acerifolia B.L. Turner, Salvia decora Epling, Salvia fusca Epling and Salvia subhastata Epling) and Coalcomán (Salvia cyanantha Epling) and two (Salvia atropaenulata Epling and Salvia filifolia Ramamoorthy) from the northeast, in Zitácuaro and Ocampo Municipalities.

Potential distribution

Individual models for *Salvia* species indicate that 59% (34,784.6 km²) of the State's total area (58,836.95 km²) is environmentally suitable to harbour its species. The species with

the largest and smallest distribution areas were *Salvia clinopodioides* Kunth with 19,895.3 km² and *S. madrigalii* with only 16.8 km² (Table 3).

The known richness in Michoacán, depicted in 15' x 15' grids (Fig. 2A), matches the potential richness (the stacking of 42 ENMs, Fig. 2B). The cutoff point for the richness map was determined by the lowest percentage of omission errors (0.7%), obtained using all 1,925 records of the 66 Salvia species here considered. Below this threshold, the suitable conditions for Salvia species cover an area of 23,541 km² (Fig. 2), 72% in the Trans-Mexican Volcanic Belt portion of the state.

Niche models are here reported for four *Salvia* species endemic to Mexico inhabiting Michoacán (Fig. 3); *Salvia albocaerulea* Linden (Fig. 3A) and *Salvia protracta* Benth (Fig. 3B) restricted to the TMVB and the SMS, respectively. *Salvia plurispicata* (Fig. 3C) and *Salvia dichlamys* Epling with wide distribution thoughout the State, occurring in the five ecoregions (Fig. 3D).

The altitude for these species is variable, although 81% of them are distributed between 2000 and 2500 m a.s.l. The number of species decreases, both at lower (< 500 m) and higher altitudes (> 3000 m).

Ecological niche models

A total of 26 uncorrelated variables (of the initial 58) were considered to run the ENMs for each *Salvia* species with sufficient (> 5) records, including eight climatic variables, seven topographic, eight edaphic and three from remote sensing data (Table 2). The variable of highest contribution value (42% of the models) was the Annual mean temperature (bio01) based on the Jackknife test provided by Maxent, with contribution percentages between 36 and 77.5%. The second most important variable was the Magnesium content (mexmg), important in 14% of the *Salvia* models, with contribution percentages from 19 to 69% and the least informative variable was the Precipitation of the coldest quarter (bio19) with 0.5%.

We obtained 42 ENMs of *Salvia* species in Michoacán (Suppl. material 3) out of 66 species, based on 404 spatially-uncorrelated records (Table 3); 10 of these models are considered exploratory having less than the five records required to be validated statistically. In addition, only three models were run using the regularisation values and the feature class obtained from the evaluation with ENMeval (Suppl. material 4); all the other models were obtained using the default parameters as specified in Maxent.

All models (made by default setting and configured) showed good performance, AUC values above 0.84 were considered good, 19% (AUC > 0.8) and 81% excellent (AUC > 0.9, Table 3). The validation of the models with the binomial test indicated that 42% models had statistically-significant prediction capacity (0.53-0.93, p-value = 0.03-3.46 x 10⁻¹⁴ respectively), while 58% of the models were not statistically significant. In contrast, the Partial-ROC tests were statistically significant (p < 0.05) for all species, with values between 1.13 and 1.99 (mean 1.58, Table 3).

Conservation status

The geographic area obtained with the assembly of models for the *Salvia* species has been profoundly affected by land use change. It has been estimated that 22% of the area has been lost in the last 20 years (1997 - 2017); TMVB ecoregion is the most severely affected, with a reduction of about 43% loss of its primary vegetation. The least affected primary vegetation loss ecoregions are JDF and the SMS (less than 10%).

Ten Protected areas in Michoacán harbour potential distribution areas of *Salvia* species. Overall, these PAs include 3.3% of the potential distribution predicted. Eight of these PAs are located in the TMVB ecoregion; unfortunately, the SMS ecoregion does not register any PA. A total of 55% *Salvia* species in the State are represented in the PAs; of these, 25 species are distributed in the Monarch Butterfly Reserve (the Monarch Butterfly Hibernation Protection Zone). Only 6% of the rare species include populations within the State's PAs.

Discussion

Species and occurrence data

The Salvia's database of Michoacán State includes more than 3,000 records, reduced to 1,925 unique records after eliminating duplicates. Although the database includes a significant number of records for the State, only 39.6% are cited in the referred floristic-taxonomic studies of relevance to the State. The two projects, focused on Salvia in Michoacán (Cornejo-Tenorio and Ibarra-Manríquez 2011, Lara-Cabrera et al. 2016), combine up to 701 specimens representing 36.4% of the total database; 685 specimens were collected before 2010 and are currently already included in online databases. Both studies cited a total of 217 reviewed specimens, representing 44% and 31.2%, respectively of those here reviewed. Although they do not explicitly indicate whether the number of specimens cited represent a total sample or a sample of the reviewed specimens, only 39.6% of the total collections for Salvia are cited.

Undoubtedly, 1,925 records representing 66 of the 69 species occurring in the State (Martínez-Gordillo et al. 2017), are a good sample of the State's generic diversity, with broad habitat diversity where the genus thrives. Many additional records here documented were reviewed by the group's specialists, although they were not cited in their floristic-taxonomic works. The 1,925 unique records included here are a summary of more than 3,000 specimens kept in about 20 different herbaria, both in Mexico and abroad (Table 1). The National Herbarium (MEXU) at the Instituto de Biología, Universidad Nacional Autónoma de México stands out, housing 64.5% of the total records (1,251); 1,159 of the 1,836 records used for the elaboration and validation of ENMs were at MEXU, constituting the main supply of information and headquarters of the study carried out here.

Patterns of distribution of Salvia in Michoacán

Our results indicate that *Salvia* species in Michoacán preferentially occur in temperate or seasonally-dry forests, the predominating biomes in the State (Gopar-Merino and Velázquez 2016); that is why more than 50% of the surface of the State is suitable for the species. Our results agree with those reported by Lara-Cabrera et al. (2016), which document higher species richness in the TMVB, where coniferous, oak and humid mountain forests abound. Similarly, Cornejo-Tenorio and Ibarra-Manríquez (2011) report that 40 *Salvia* species are distributed in the pine-oak and oak forests. Thus, our ENM are consistent with the previously reported environmental preferences for the genus.

The majority of *Salvia* species act as generalists to environmental conditions throughout their distribution in Michoacán. Only temperature had the highest variable contribution amongst all biomes; temperature is frequently documented as the dominant abiotic driver in determining plant distributions (Körner and Hiltbrunner 2018, Scherrer and Guisan 2019). Mean temperature is associated with plant growth, acting mainly in a gradual manner (Körner et al. 2016), determining whether a plant species will be present (Guisan et al. 1998), particularly in the mountainous landscape, such as is characteristic in the TMVB (Cornejo-Tenorio and Ibarra-Manríquez 2011, Lara-Cabrera et al. 2016). Oddly, precipitation was not a significant variable; a correlation between precipitation and species richness at regional scale in these environments has been well documented, the greater the precipitation, the greater the species richness (Pau et al. 2012). However, precipitation's contribution here is low, probably due to scarce records from the Balsas Dry Forests and JDF (Fig. 2), where rainfall is below 1,000 mm.

Rare species (< 3 records), although not modelled, show a preference for similar environmental conditions as compared to modelled species or widely-distributed species (> 3 records). For example, the mean temperature of rare species (18°C) is slightly higher than that of widely-distributed species (15.7°C), while in precipitation, the mean values are very similar, 1,057 mm for rare species and 1,058 mm for widely-distributed species. Furthermore, several state endemic *Salvia* species require very specific conditions; *S. madrigalii*, *S. synodonta* and *S. cyanantha*, are known only from two sites with practically the same climatic characteristics. The specificity of their habitat, as a result of their restricted distribution, makes these species particularly susceptible to extinction (Isik 2011). Despite having information on the optimal environmental conditions of this species group, very little is known about the biological or physiological aspects that regulate it.

Environmental suitability is indicated for the genus along the SMS (Fig. 2); however, further fieldwork for the region could add important floristic information (Cornejo-Tenorio and Ibarra-Manríquez 2011, Lara-Cabrera et al. 2016), as far as the optimal environmental conditions where they thrive. The above is a particular way for endemics or rare species that would allow us to confirm or improve the distribution patterns reported in this study.

Cornejo-Tenorio and Ibarra-Manríquez (2011) mention altitude as an important variable in the distribution from *Salvia*. The inclusion of this variable in the potential geographic range

of *Salvia* in Michoacán confirms its importance in explaining the species distribution. The genus inhabits altitudes from 200 to 3,600 m, as reported by Lara-Cabrera et al. (2016); however, species richness is not evently distributed throughout this altitudinal range, with middle altitudes (2,000-2,500 m) as the richest in species (52 species) and decreasing richness both at lower and higher altitudes. These data coincide with those published by Cornejo-Tenorio and Ibarra-Manríquez (2011), who mention that the highest richness was found above 2,000 m altitude (35 species).

Although the large amount of information in the databases would seem sufficient for ENMs and thus inferring many species distribution, this was not always the case. When performing statistical analyses to develop more robust and precise models, such as eliminating spatial autocorrelation of points to reduce sampling bias or to optimise the specific configuration for each model adjusted for each species data (Boria et al. 2014, Radosavljevic and Anderson 2014, Warren et al. 2014), it became clear that we did not have enough independent records. In the case of Salvia in Michoacán, despite several works specifically focused on these species (Cornejo-Tenorio and Ibarra-Manríquez 2011, Lara-Cabrera et al. 2016, Zamudio and Bedolla-García 2018), we still did not have sufficient data for most species to develop ENMs under specific configurations. We found that 78% of the purged data for Salvia are spatially correlated, which is likely the result of sampling bias to areas of easy access or of particular interest, resulting in several records from a single site for the same species. Due to this strong point correlation, we were able to obtain only three configured models (models whose rigour was increased by adjusting their configuration or balancing their complexity). We corroborate the guidelines by other authors, at least 20 records are necessary to obtain and reach the predictive capacity for models (Stockwell and Peterson 2002, Mateo et al. 2010, van Proosdij et al. 2016). In our case, using few records to obtain regularisation values, we obtained models with low predictive value, resulting in high rates of model overprediction and AUC values below 0.5. For this reason, we too recommended that the tests to obtain regularisation values and make configured models be used only when more than 20 records are available; the larger the number of records, the better the model performance (Stockwell and Peterson 2002) as shown in Salvia.

The AUC has been criticised as a tool to evaluate only the performance of presence models, because of its dependence on prevalence; therefore, it is not considered a precise performance index (Lobo et al. 2008, van Proosdij et al. 2016). For this reason, other performance evaluation metrics, such as Partial ROC for continuous data (Peterson and Nakazawa 2008) and the binomial test for binary models (Anderson et al. 2002), were implemented. We included these test results as additional evaluation measures (Table 3). The number of non-significant binomial tests (58% of the models) correspond to the number of species with less than five records. Blair et al. (2013) obtained similar results when using ten records; below this sample size, their binomial tests were not significant (p > 0.05). We recommend reaching a minimum number of 20 independent points of occurrence to construct models with better predictive power that pass the test of sensitivity to small samples.

High deforestation rates in the north-central region of Michoacán, is the main cause of habitat loss. The most vulnerable vegetation types are temperate forests, constantly being replaced by avocado orchards (Mas et al. 2017) amongst other threats, thus decreasing the original distribution of many species, as we found here for *Salvia*.

Implications for conservation

Protected areas represent a key strategy for biodiversity preservation (Hannah et al. 2007, Belote and Wilson 2020); however, most are limited to small geographic areas that do not represent the region's diversity for *Salvia*. Only 3% of its potential distribution is present in PAs, comprising 55% of its species richness. In order to increase the representation of these species within the PAs, it has been proposed to raise the conservation status to larger ecosystems; this approach would allow us to increase the representation of vegetation types and with it, the number of conserved species (Belote and Wilson 2020).

The results point to the TMVB, the SMS and the JDF as priority ecoregions for *Salvia* conservation in Michoacán, where the most species richness and rare species (the only known locality for some of the species) are located. Despite the high species richness, few PAs in the State have been thoroughly sampled (Cornejo-Tenorio and Ibarra-Manríquez 2011). The State PAs (Fig. 2) correspond with only 3% of *Salvia*'s potential distribution area and none of them includes sites for rare species. For example, in the south-western part of the State, no PA is recorded, although nine rare species are recorded in this region and three of them are only known from one locality. Such is the case of *S. acerifolia* (JDF) or *S. cyanantha* and *S. subhastata* (both from the SMS); conservation strategies should be focus on these sites. In the case of the TMVB, seven rare species are recorded and only eight PAs have been decreed; thus, future conservation efforts should aim at their expansion, to include a greater number of species and thereby conserve rare species. This ecoregion is suffering the greatest natural habitats loss (56%). The areas of highest richness identified here (darker areas of Fig. 2) could be used to propose specific sites for the expansion of protected areas.

Acknowledgements

We are grateful to CONABIO and the Instituto de Biología, UNAM, for access to the information stored in the SNIB-REMIB and UNIBIO databases, respectively, which constituted a fundamental part of the analyses presented here. Thanks to Enrique Ortiz and Joselin Cadena Rodriguez for reviewing a draft of the manuscript and two anonymous reviewers; their assistance improved substantially the document. In addition, Enrique Ortiz helped in several facets of the study with his technical assistance. We appreciate Lynna Kiere's editing of the English version.

Hosting institution

National Autonomous University of Mexico, Mexico, City.

Author contributions

M.F.T. and J.L.V., conceived the project, designed the methods, conducted the statistical analysis and species distribution modelling. M.F.T., J.L.V and S.I.L.C interpreted the results, authored the writing and approved the final draft.

Conflicts of interest

The authors declare that there is no conflict of interest regarding the publication of this manuscript.

References

- Aiello-Lammens ME, Boria RA, Radosavljevic A, Vilela B, Anderson RP (2015) spThin: An R package for spatial thinning of species occurrence records for use in ecological niche models. Ecography 38: 541-545. https://doi.org/10.1111/ecog.01132
- Akaberi M, Iranshahi M, Mehri S (2016) Molecular signaling pathways behind the biological effects of Salvia species diterpenes in neuropharmacology and cardiology. Phytotherapy Research 30: 878-893. https://doi.org/10.1002/ptr.5599
- Anderson RP, Gmez-Laverde M, Peterson AT (2002) Geographical distributions of spiny pocket mice in South America: insights from predictive models. Global Ecology Biogeography 11: 131-141. https://doi.org/10.1046/j.1466-822X.2002.00275.x
- Argumedo DR, Parra-Delgado H, Ramírez AM, Nieto CA, Martnez-Vzquez M (2003)
 Isolation and chemical transformations of some anti-inflammatory triterpenes from Salvia mexicana L. var. minor Benth. Revista de la Sociedad Qumica de Mexico 47: 167-172.
- Bedolla-García B, Lara-Cabrera S, Zamudio S (2011) Dos nuevas especies de Salvia (Lamiaceae) del centro occidente de México. Acta Botanica Mexicana 95: 51-63. https://doi.org/10.21829/abm95.2011.267
- Belote RT, Wilson MB (2020) Delineating greater ecosystems around protected areas to guide conservation. Conservation Science and Practice 2 (6): e196. https://doi.org/10.1111/csp2.196
- Blair ME, Sterling EJ, Dusch M, Raxworthy CJ, Pearson RG (2013) Ecological divergence and speciation between lemur (Eulemur) sister species in Madagascar. Journal of Evolutionary Biology 26: 1790-1801. https://doi.org/10.1111/jeb.12179
- Boria RA, Olson LE, Goodman SM, Anderson RP (2014) Spatial filtering to reduce sampling bias can improve the performance of ecological niche models. Ecological Modelling 275: 73-77. https://doi.org/10.1016/j.ecolmodel.2013.12.012

- Caballero J, Cortés L (2001) Plantas, cultura y sociedad. Estudio sobre la relación entre seres humanos y plantas en los albores del siglo XXI. Universidad Autónoma Metropolitana, Unidad Iztapalapa y Secretara del Medio Ambiente, Recursos Naturales y Pesca, México, D.F., 317 pp. [ISBN 970-654-782-7]
- Cahill JP (2003) Ethnobotany of Chia, Salvia hispanica L. (Lamiaceae). Economic Botany 57: 604-618. https://doi.org/
 10.1663/0013-0001(2003)057[0604:EOCSHL]2.0.CO;2
- Castillo M, Michán L, Martínez AL (2014) La biocuración en biodiversidad: proceso, aciertos, errores, soluciones y perspectivas. Acta Botanica Mexicana 108: 81-103. https://doi.org/10.21829/abm108.2014.199
- Chapman AD (2005) Principles and Methods of Data Cleaning Primary Species and Species Occurrence Data, version 1.0. Report for the Global Biodiversity Information Facility, Copenhagen.
- CONANP (2010) Áreas Naturales Protegidas Federales de México. http://www.conanp.gob.mx/sig/informacion/info.htm. Accessed on: 2020-3-26.
- Cornejo-Tenorio G, Ibarra-Manríquez G (2011) Diversidad y distribución del género Salvia (Lamiaceae) en Michoacán, México. Revista Mexicana Biodiversidad 82: 1279-1296.
- Cruz-Cárdenas G, López-Mata L, Villaseñor JL, Ortíz E (2014a) Potential species distribution modeling and the use of principal component analysis as predictor variables. Revista Mexicana de Biodiversidad 85: 189-199. https://doi.org/10.7550/rmb.36723
- Cruz-Cárdenas G, López-Mata L, Ortíz-Solorio CA, Villaseñor JL, Ortíz E, Teodoro SJ, Estrada-Godoy F (2014b) Interpolation of Mexican soil properties at a scale of 1:1,000,000. Geoderma 213: 29-35. https://doi.org/10.1016/j.geoderma.2013.07.014
- Elith J, Phillips S, Hastie T, Dudik M, Chee Y, Yates CA (2011) A statistical explanation
 of MaxEnt for ecologists. Diversity and Distributions 17: 43-57. https://doi.org/10.1111/j.1472-4642.2010.00725.x
- Espejo-Serna A, Ramamoorthy TP (1993) Revisión taxonómica de Salvia sección Sigmoideae (Lamiaceae). Acta Botanica Mexicana 23: 65-102. https://doi.org/10.21829/abm23.1993.675
- ESRI (2013) ArcGIS: Environmental Systems Research Institute. https://www.esri.com/es-es/home
- Fourcade Y, Besnard AG, Secondi J (2018) Paintings predict the distribution of species, or the challenge of selecting environmental predictors and evaluation statistics. Global Ecology and Biogeography 27: 245-256. https://doi.org/10.1111/geb.12684
- González-Chávez MM, Alonso-Castro AJ, Zapata-Morales JR, Arana-Argañez V, Torres-Romero JC, Medina-Rivera YE, Sánchez-Mendoza E, Perez-Gutierrez S (2018) Anti-inflammatory and antinociceptive effects of tilifodiolide, isolated from Salvia tiliifolia Vahl (Lamiaceae). Drug Development Research 9: 165-172. https://doi.org/10.1002/ddr.21432
- González-Gallegos JG (2014) Revision of Salvia subg. Calosphace sect. Membranaceae (Lamiaceae). Telopea 16: 43-81. https://doi.org/10.7751/telopea20147483
- González-Gallegos JG, Castro-Castro A, Quintero-Fuentes V, Mendoza-López ME, De Castro-Arce E (2016) Revisión taxonómica de Lamiaceae del occidente de México. Ibugana 7: 3-345.

- Gopar-Merino LF, Velázquez A (2016) Componentes del paisaje como predictores de cubiertas de vegetación: estudio de caso del estado de Michoacán, México.
 Investigaciones Geográficas, Boletin 90: 75-88. https://doi.org/10.14350/rig.46688
- Guisan A, Theurillat J-, Kienast F (1998) Predicting the potential distribution of plant species in an alpine environment. Journal of Vegetation Science 9: 65-74. https://doi.org/10.2307/3237224
- Hannah L, Midgley G, Andelman S, Araújo M, Hughes G, Martinez-Meyer E, Pearson R, Williams P (2007) Protected area needs in a changing climate. Frontiers in Ecology and the Environment 5: 131-138. https://doi.org/10.1890/1540-9295(2007)5[131:PANIAC]2.0.CO:2
- Hernandez PA, Graham CH, Master LL, Albert DL (2006) The effect of simple size and species characteristics on performance of different species distribution modeling methods. Ecography 29: 771-785. https://doi.org/10.1111/j.0906-7590.2006.04700.x
- 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. https://doi.org/10.1002/joc.1276
- Hijmans RJ, Graham CH (2006) The ability of climate envelope models to predict the effect of climate change on species distributions. Global Change Biology 12: 2272-2281. https://doi.org/10.1111/j.1365-2486.2006.01256.x
- Hijmans RJ, Phillips S,, Leathwick J, Elith J (2017) dismo: Species distribution modeling. R package. 1.1-4. URL: http://cran.r-project.org/package=dismo
- Iltis HH, González-Gallegos JG, Cochrane TS, Vázquez-García JA (2012) A new species and a new subspecies of Salvia (Lamiaceae) from Jalisco and Michoacán, Mexico. Brittonia 64 (4): 343-352. https://doi.org/10.1007/s12228-012-9237-1
- INEGI (1997) Conjunto de datos vectoriales de la carta de Uso del suelo y vegetación. https://www.inegi.org.mx/app/mapas/. Accessed on: 2020-3-29.
- INEGI (2017) Conjunto de datos vectoriales de la carta de Uso del suelo y vegetación. Escala 1:250 000. Serie VI. https://www.inegi.org.mx/app/mapas/. Accessed on: 2020-3-29.
- Isik K (2011) Rare and endemic species: why are they prone to extinction? Turkish Journal of Botany 35: 411-417. https://doi.org/10.3906/bot-1012-90
- Jetz W, McGeoch MA, Guralnick R, Ferrier S, Beck J, Costello MJ, Turak E (2019)
 Essential biodiversity variables for mapping and monitoring species populations. Nature
 Ecology and Evolution 3: 539-551. https://doi.org/10.1038/s41559-019-0826-1
- Kass J, Vilela B, Aiello-Lammens M, Muscarella R, Merow C, Anderson RP (2018)
 Wallace: a flexible platform for reproducible modeling of species niches and distributions built for community expansion. Methods in Ecology and Evolution 9: 1151-1156. https://doi.org/10.1111/2041-210X.12945
- Körner C, Basler D, Hoch G, Kollas C, Lenz A, Randin CF, Vitasse Y, Zimmermann NE, Turnbull M (2016) Where, why and how? Explaining the lowtemperature range limits of temperate tree species. Journal of Ecology 104: 1076-1088. https://doi.org/10.1111/1365-2745.12574
- Körner C, Hiltbrunner E (2018) The 90 ways to describe plant temperature.
 Perspectives in Plant Ecology, Evolution and Systematics 30: 16-21. https://doi.org/10.1016/j.ppees.2017.04.004

- Lara-Cabrera SI, Bedolla-García BY, Zamudio S, Domínguez-Vázquez G (2016)
 Diversidad de Lamiaceae en el estado de Michoacán, México. Acta Botanica Mexicana
 116: 107-149. https://doi.org/10.21829/abm116.2016.1120
- Liu C, Newell G, White M (2016) On the selection of thresholds for predicting species occurrence with presence-only data. Ecology and Evolution 6 (1): 337-348. https://doi.org/10.1002/ece3.1878
- Li Y, Wen J, Ren Y, Zhang J (2019) From seven to three: Integrative species
 delimitation supports major reduction in species number in *Rhodiola* section *Trifida*(Crassulaceae) on the Qinghai-Tibetan Plateau. TAXON 68 (2): 268-279. https://doi.org/10.1002/tax.12052
- Lobo JM, Jiménez-Valverde A, Real R (2008) AUC: a misleading measure of the performance of predictive distribution models. Global Ecology and Biogeography 17: 145-151. https://doi.org/10.1111/j.1466-8238.2007.00358.x
- Maqueda AE, Valle M, Addy PH, Antonijoan RM, Puntes M, Coimbra J, Ballester MR, Garrido M, Gonzlez M, Claramunt J, Barker S, Johnson MW, Griffiths RR, Riba J (2015) Salvinorin-A induces intense dissociative effects, blocking external sensory perception and modulation interoception and sense of body ownership in humans. International Journal of Neuropsychopharmacology 18: 1-14. https://doi.org/10.1093/ijnp/pyv065
- Martínez-Gordillo M, Fragoso-Martínez I, García-Peña MR, Montiel O (2013) Géneros de Lamiaceae de México, diversidad y endemismo. Revista Mexicana Biodiversidad 84: 30-86. https://doi.org/10.7550/rmb.30158
- Martínez-Gordillo M, Bedolla-García B, Cornejo-Tenorio G, Fragoso-Martínez I, García-Peña MR, González-Gallegos JG, Lara-Cabrera SI, Zamudio S (2017) Lamiaceae de México. Botanical Sciences 95: 780-806. https://doi.org/10.17129/botsci.1871
- Mas JF, Lemoine-Rodríguez R, González R, López-Sánchez J, Piña-Garduño A, Herrera-Flores E (2017) Evaluación de las tasas de deforestación en Michoacán a escala detallada mediante un método híbrido de clasificación de imágenes SPOT. Madera y Bosques 23 (2): 119-132. https://doi.org/10.21829/myb.2017.2321472.
- Mateo RG, Felicísimo AM, Muñoz J (2010) Effects of the number of presences on reliability and stability of MARS species distribution models: the importance of regional niche variation and ecological heterogeneity. Journal of Vegetation Science 21: 908-922. https://doi.org/10.1111/j.1654-1103.2010.01198.x
- Moss L, Rouse M, Wesnes AK, Moss M (2010) Differential effects of the aromas of Salvia species on memory and mood. Human Psychopharmacology: Clinical and Experimental 25: 388-396. https://doi.org/10.1002/hup.1129
- Munguía-Lino G, Vázquez-García LM, López-Sandoval JA (2010) Plantas silvestres ornamentales comercializadas en los mercados de La Flor de Tenancingo y Jamaica, México. Polibotánica 29: 281-308.
- Muscarella R, Galante PJ, Soley-Guardia M, Boria RA, Kass JM, Uriarte M, Anderson RP (2014) ENMeval: An R package for conducting spatially independent evaluations and estimating optimal model complexity for Maxent ecological niche models. Methods in Ecology and Evolution 5: 1198-1205. https://doi.org/10.1111/2041-210X.12261
- Olson DM, Dinerstein E, Wikramanayake ED, Burgess ND, Powell GV, Underwood EC, Damico JA, Itoua I, Strand HE, Morrison JC, Loucks CJ, Allnutt TE, Rickett TH, Kura Y, Lamoreux JF, Wettendgel WW, Hedao P, Kassem KR (2001) Terrestrial ecoregions of the world: a new map of life on Earth. Bioscience 51 (11): 933-938. https://doi.org/10.1641/0006-3568(2001)051[0933:TEOTWA]2.0.CO;2

- Olvera-Mendoza EI, Bedolla-García BY, Lara-Cabrera SI (2017) Revisión taxonómica de Salvia subgénero Calosphace sección Scorodoniae (Lamiaceae), endémica de México. Acta Botanica Mexicana 118: 7-40. https://doi.org/10.21829/abm118.2017.1198
- Osorio-Olvera L, Barve V, Barve N, Soberón J, Falconi M (2018) ntbox: From getting biodiversity data to evaluating species distribution models in a friendly GUI environment. R package. 0.2.5.4. URL: https://github.com/luismurao/ntbox"
- Pau S, Gillespie TW, Wolkovich EM (2012) Dissecting NDVI-species richness relationships in Hawaiian dry forests. Journal of Biogeography 39 (9): 1678-1686. https://doi.org/10.1111/j.1365-2699.2012.02731.x
- Peterson AT, Nakazawa Y (2008) Environmental data sets matter in ecological niche modelling: an example with Solenopsis invicta and Solenopsis richteri. Global Ecology and Biogeography 17: 135-144. https://doi.org/10.1111/j.1466-8238.2007.00347.x
- Peterson AT, Soberón J, Pearson RG, Anderson RP, Martínez-Meyer E, Nakamura M, Araújo MB (2011) Ecological niches and geographic distributions (MPB-49). Princeton University Press, New Jersey, USA. https://doi.org/10.23943/princeton/9780691136868.001.0001
- Phillips SJ, Anderson RP, Schapire RE (2006) Maximum entropy modeling of species geographic distributions. Ecological Modelling 190: 231-259. https://doi.org/10.1016/j.ecolmodel.2005.03.026
- Phillips SJ, Dudík M, Schapire RE (2017) Maxent software for modeling species niches and distributions. 3.4.1. URL: https://biodiversityinformatics.amnh.org/open_source/maxent/
- Ponder WF, Carter GA, Flemons P, Chapman RR (2001) Evaluation of museum collection data for use in biodiversity assessment. Conservation Biology 15: 648-657. https://doi.org/10.1046/j.1523-1739.2001.015003648.x
- Radosavljevic A, Anderson RP (2014) Making better Maxent models of species distributions: Complexity, overfitting and evaluation. Journal of Biogeography 41: 629-643. https://doi.org/10.1111/jbi.12227
- Ramamoorthy TP, Elliot M (1998) Lamiaceae de México: diversidad, distribución, endemismo y evolución. In: Ramamoorthy TP, Bye R, Lot A, Fa J (Eds) Diversidad Biológica de México: Orígenes y Distribución. México. Instituto de Biología, Universidad Nacional Autónoma de México, Mexico city, 792 pp.
- R Development Core Team (2016) R: A language and environment for statistical computing. R Foundation for Statistical Computing, Vienna, Austria. 3.3.2. URL: http://www.r-project.org/
- Scherrer D, Guisan A (2019) Ecological indicator values reveal missing predictors of species distributions. Nature Scientific Reports 9: 3061. https://doi.org/10.1038/s41598-019-39133-1
- Soberón JM, Peterson AT (2005) Interpretation of models of fundamental ecological niches and species' distributional areas. Biodiversity Informatics 2: 1-10. https://doi.org/10.17161/bi.v2i0.4
- Staude IR, Navarro LM, Pereira HM (2019) Range size predicts the risk of local extinction from habitat loss. Global Ecology and Biogeography 29: 16-25. https://doi.org/10.1111/geb.13003
- Stockwell DR, Peterson AT (2002) Effects of sample size on accuracy of species distribution models. Ecological Modelling 148: 1-13. https://doi.org/10.1016/ \$0304-3800(01)00388-X

- van Proosdij AS, Sosef MS, Wieringa JJ, Raes N (2016) Minimum required number of specimen records to develop accurate species distribution models. Ecography 39: 542-552. https://doi.org/10.1111/ecog.01509
- Varela S, Anderson RP, García-Valdés R, Fernández-González F (2014) Environmental filters reduce the effects of sampling bias and improve predictions of ecological niche models. Ecography 37: 1-8. https://doi.org/10.1111/j.1600-0587.2013.00441.x
- Villaseñor JL, Téllez-Valdés O (2004) Distribución potencial de las especies del género Jefea (Asteraceae) en México. Anales del Instituto de Biología UNAM (Serie Botánica) 75: 205-220.
- Villaseñor JL (2016) Checklist of the native vascular plants of Mexico. Revista Mexicana de Biodiversidad 87 (3): 559-902. https://doi.org/10.1016/j.rmb.2016.06.017
- Walker JB, Sytsma KJ, Treutlein J, Wink M (2004) Salvia (Lamiaceae) is not monophyletic: Implications for the systematics, radiation, and ecological specializations of Salvia and tribe Mentheae. American Journal of Botany 91: 1115-1125. https://doi.org/10.3732/ajb.91.7.1115
- Warren DL, Glor RE, Turelli M (2008) Environmental niche equivalency versus conservatism: quantitative approaches to niche evolution. Evolution 62: 2868-2883. https://doi.org/10.1111/j.1558-5646.2008.00482.x
- Warren DL, Wright AN, Seifert SN, Shaffer HB (2014) Incorporating model complexity
 and spatial sampling bias into ecological niche models of climate change risks faced by
 90 California vertebrate species of concern. Diversity and Distributions 20: 334-343.
 https://doi.org/10.1111/ddi.12160
- Whittaker RJ, Araújo MB, Jepson P, Ladle RJ, Watson JE, Willis KJ (2005)
 Conservation biogeography: Assessment and prospect. Diversity and Distributions 11:
 3-23. https://doi.org/10.1111/j.1366-9516.2005.00143.x
- Zamudio A, Bedolla-García BY (2018) Salvia madrigalii (Lamiaceae), una especie nueva de Michoacán, México. Brittonia 70 (1): 76-83. https://doi.org/10.1007/s12228-017-9491-3

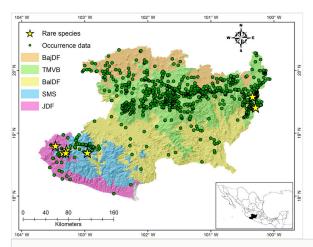


Figure 1.

Known distribution of the genus *Salvia* based on collecting effort (green points) in Michoacán. The yellow stars indicate the location of species that have only one record in Michoacán. Colours show the boundaries of each ecoregion. BajDF: Bajío Dry Forests, TMVB: Trans-Mexican Volcanic Belt, BalDF: Balsas Dry Forests, SMS: Sierra Madre del Sur, JDF: Jalisco Dry Forest

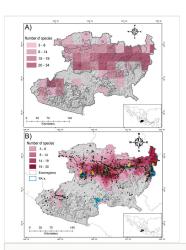


Figure 2.

Known and potential distribution of the genus *Salvia*. **A.** The State of Michoacán, Mexico, divided into 15' × 15' squares showing the known distribution of the genus *Salvia* in the State; **B.** Potential distribution of the genus *Salvia* in Michoacán (pink colour). Darker colours correspond to areas with more assemblage of *Salvia* species. The points indicate the collection sites. Blue lines show the boundaries of the Protected Areas and grey boundaries correspond to the limits of the ecoregions. 1: Zicuirán-Infiernillo, 2: Pico de Tancítaro, 3: Barranca del Cupatitzio, 4: Insurgente José María Morelos, 5: Mariposa Monarca

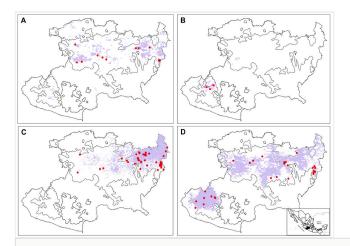


Figure 3.

Examples of potential distribution areas of species of *Salvia* endemic to Mexico (lilac colour) and representative of the Michoacán ecoregions (black boundaries within the State of Michoacán). **A.** *Salvia albocaerulea;* **B.** *Salvia protracta;* **C.** *Salvia plurispicata* (endemic to Michoacán); **D.** *Salvia dichlamys*. The red circles show the points of occurrence of the species.

Table 1.

Floristic and taxonomic studies accounting for species diversity of *Salvia* in the Mexican State of Michoacán.

Study	Species in Michoacán	Total number of records cited for Michoacán	Number of records in <i>Salvia</i> 's data base
Espejo-Serna and Ramamoorthy (1993)	4	52	52
Cornejo-Tenorio and Ibarra-Manríquez (2011)	64	493	439
González-Gallegos (2014)	3	5	1
González-Gallegos et al. (2016)	14	189	97
Lara-Cabrera et al. (2016)	62	695	479
Martínez-Gordillo et al. (2017)	69	None	None
Olvera-Mendoza et al. (2017)	3	19	6
Salvia's data base (this study)	70	768	1,925

Table 2.

Variables used to estimate the ecological niche models of *Salvia* species in Michoacán. The variables with the highest and lowest contribution in the ENMS of *Salvia* species are highlighted in hold

Туре	Variable
Climatic	bio01 (Annual mean temperature)
	bio02 (Average daytime variation)
	bio03 (Isothermality)
	bio14 (Precipitation of driest month)
	bio15 (Seasonality of precipitation)
	bio18 (Precipitation of the warmest quarter)
	bio19 (Precipitation of the coldest quarter)
	evaanual (Annual real evapotranspiration)
Topographic	aspect (Orientation 0° to 90°)
	convrgin (Convergence index)
	dah (Diurnal anisotropic heating)
	mexslope (Slope)
	runoff (Flow)
	twi (Topographic moisture index)
	vrm (Vector rugosity measure)
Edaphic	mexca (Calcium)
	mexce (Electrical conductivity)
	mexco (Organic carbon)
	mexk (Potassium)
	mexmg (Magnesium)
	mexmo (Organic material)
	mexna (Sodium)
	mexras (Sodium absorption radius)
*MODIS	modismar (Normalised index of vegetation March)
	modissep (Normalised index of vegetation September)
	hummodis2009 (Normalised index of vegetation humid months)

Table 3.

Species of Salvia in Michoacán for which ecological niche models could be constructed. The number of records used to build the models and the surface estimated by the models are indicated. The values of the AUC and Partial-ROC tests are indicated for each species. Bold type indicates species that did not pass the binomial test (p > 0.05). The exploratory models are marked with an asterisk.

Species	Total records/ records to build the models	Potential area (km²)	AUC	Partial ROC	Binomial test
Salvia albocaerulea Linden	14/5	3286.3	0.972	1.7	0.193 (0.64)
Salvia amarissima Ortega	15/7	8281.8	0.934	1.90	0.050 (0.49)
*Salvia assurgens Kunth	44/4	3727.1	0.962	1.38	0.183 (1)
*Salvia carnea Kunth	6/4	1576.0	0.986	1.83	0.355 (0.29)
Salvia clinopodioides Kunth	47/12	19895.3	0.837	1.56	0.913 (1.16 x 10 ⁻¹⁰)
Salvia curviflora Benth.	8/5	3388.1	0.971	1.91	0.473 (0.06)
Salvia dichlamys Epling	32/15	11831.4	0.89	1.29	0.260 (0.69)
Salvia elegans Vahl	130/25	11912.2	0.874	1.56	0.795 (3.46 x 10 ⁻¹³)
Salvia fulgens Cav.	95/10	9306.3	0.909	1.67	0.770 (1.36 x 10 ⁻¹¹)
*Salvia gravida Epling	8/3	1982.4	0.988	1.97	0.607 (0.02)
Salvia gesneriiflora Lindl. & Paxton	57/17	3516.7	0.985	1.59	0.575 (0.006)
Salvia helianthemifolia Benth.	48/7	2873.0	0.979	1.93	0.930 (4.55 x 10 ⁻¹³)
*Salvia hispanica L.	23/4	1853.2	0.974	1.23	0.003 (1)
Salvia iodantha Fernald	138/20	7989.4	0.896	1.49	0.556 (0.002)
Salvia laevis Benth.	57/18	12436.1	0.932	1.59	0.887 (3.73 x 10 ⁻¹¹)
*Salvia languidula Epling	9/4	291.8	0.995	1.85	0.009 (0.98)
Salvia lasiocephala Hook. & Arn.	34/14	8340.8	0.934	1.17	0.075 (1)
Salvia lavanduloides Kunth	102/15	16136.8	0.883	1.25	0.647 (6.33 x 10 ⁻⁶)
Salvia leptostachys Benth.	16/5	2505.2	0.987	1.46	0.063 (0.89)
Salvia longispicata M. Martens & Galeotti	76/13	12802.5	0.928	1.41	0.578 (0.003)
Salvia longistyla Benth.	37/8	18161.9	0.926	1.45	0.712 (5.19 x 10 ⁻⁵
*Salvia madrigalii Zamudio & Bedolla	4/3	16.8	1	1.99	0.473 (0.06)
Salvia melissodora Lag.	11/5	2705.5	0.972	1.61	0.000 (1)
Salvia mexicana L.	206/24	7161.9	0.861	1.51	0.508 (0.03)

Salvia microphylla Kunth	38/6	12920.7	0.9	1.31	0.521 (0.03)
Salvia misella Kunth	55/19	10142.5	0.854	1.23	0.258 (0.84)
Salvia mocinoi Benth.	35/14	10954.4	0.925	1.71	0.793 (1.05 x 10 ⁻⁵
Salvia patens Cav.	20/12	2008.3	0.981	1.94	0.529 (0.04)
Salvia plurispicata Epling	55/13	6165.0	0.963	1.69	0.663 (0.0003)
Salvia polystachia Cav.	96/11	18053.7	0.838	1.39	0.768 (2.31 x 10 ⁻¹¹)
*Salvia protracta Benth.	4/3	171.0	0.997	1.99	0.0 (1)
Salvia prunelloides Kunth	13/5	707.6	0.993	1.71	0.129 (0.77)
Salvia purpurea Cav.	62/10	9902.8	0.899	1.32	0.435 (0.24)
*Salvia reflexa Hornem.	4/4	5217.0	0.948	1.87	0.0 (1)
Salvia reptans Jacq.	55/12	7664.7	0.952	1.47	0.577 (0.004)
*Salvia rhyacophila (Fernald) Epling	6/4	2073.6	0.985	1.95	0.0 (1)
Salvia sessei Benth.	34/6	16430.3	0.877	1.32	0.282 (0.73)
*Salvia setulosa Fernald	6/4	912.5	0.99	1.97	0.0 (1)
*Salvia stachyoides Kunth	11/4	2404.5	0.976	1.66	0.200 (0.73)
Salvia thyrsiflora Benth.	70/15	6612.8	0.882	1.13	0.195 (1)
Salvia tiliifolia Vahl	40/7	12192.5	0.913	1.25	0.282 (0.73)
Salvia uruapana Fernald	15/8	7186.5	0.933	1.21	0.076 (0.81)

Supplementary materials

Suppl. material 1: List of specimens of *Salvia* species occurring in the State of Michoacán, Mexico considered in this study.

Authors: Mayra Flores-Tolentino, José Luis Villaseñor and Sabina Lara-Cabrera

Data type: Specimens

Brief description: Each record is made up of the collector's last name and collecting number; in parentheses the herbaria where the specimen is stored, followed by the coordinates in decimal degrees. In brackets, the publications in which the records are cited is indicated. 1: Espejo-Serna and Ramamoorthy (1993); 2: Cornejo-Tenorio and Ibarra Manríquez (2011); 3: Iltis et al. (2012); 4: González-Gallegos et al. (2014); 5: González-Gallegos et al. (2016); 6: Lara-Cabrera et al. (2016);

7: recorded in SNIB-REMIB and/or MEXU-UNIBIO databases.

Download file (26.68 kb)

Suppl. material 2: Species of *Salvia* recorded in the State of Michoacán and considered in this study

Authors: Mayra Flores-Tolentino, José Luis Villaseñor and Sabina Lara-Cabrera

Data type: List of species

Brief description: The number of records obtained for each species after cleaning the data from

the SNIB-REMIB and MEXU-UNIBIO databases are indicated.

Download file (4.69 MB)

Suppl. material 3: Ecological niche models of 42 Salvia species.

Authors: Mayra Flores-Tolentino and José Luis Villaseñor

Data type: Images

Brief description: Ecological niche models of 42 Salvia species. The points on each map show

the collecting localities.

Download file (4.69 MB)

Suppl. material 4: Configuration of Maxent

Authors: Mayra Flores-Tolentino and José Luis Villaseñor

Data type: Evaluation metrics

Brief description: Evaluation metrics of Maxent ENMs generated with the ENMeval programme for the three species of *Salvia* for which the statistical validation could be carried out. FC= Feature

class (Logistic); MR= Multiple regularisation; ΔAICc= Delta Akaike Information Criterion.

Download file (33.00 kb)