Centre-periphery approaches based on geography, ecology and historical climate stability: what explains the variation in morphological traits of

Bulnesia sarmientoi?

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- **Background and Aims** The Centre–Periphery Hypothesis posits that higher species performance is expected in geographic and ecological centres rather than in peripheral populations. However, this is not the commonly found pattern; therefore, alternative approaches, including the historical dimension of species geographical ranges should be explored. Morphological functional traits are fundamental determinants of species performance, commonly related to environmental stability and productivity. We tested whether or not historical processes may have shaped variations in tree and leaf traits of the Chaco tree *Bulnesia sarmientoi*.
- Methods Morphological variation patterns were analysed from three centre-periphery approaches: geographical, ecological and historical. Tree (stem and canopy) and leaf (leaf size and specific leaf area) traits were measured in 24 populations across the species range. A Principal Component Analysis was performed on morphological traits to obtain synthetic variables. Linear Mixed-Effects Models were used to test which of the implemented centre-periphery approaches significantly explained trait spatial patterns.
- **Key Results** The patterns retrieved from the three centre-periphery approaches were not concordant. The historical approach revealed that trees were shorter in center populations than in the periphery. Significant differences in leaf traits were observed between the geographical centre and the periphery, mainly due to low specific leaf area values towards the geographical centre. We did not find any pattern associated with the ecological centre-periphery approach.
- **Conclusions** The decoupled response between leaf and tree traits suggests that these sets of traits respond differently to processes occurring at different times. The geographical and historical approaches showed centres with extreme environments in relation to their respective peripheries, but the historical centre has also been a

climatically stable area since the Last Glacial Maximum. The historical approach allowed for the recovery of historical processes underlying tree traits variation, highlighting that centre-periphery delimitations should be based on a multi-approach framework.

Key words: Centre-Periphery Hypothesis, Gran Chaco Americano, ecological niche modelling, geographical centre, intraspecific traits, *Bulnesia sarmientoi*, marginal plant populations, niche centroid, refugium.

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INTRODUCTION

The Centre-Periphery Hypothesis (CPH) is one of the main frameworks used to study population performance across a species range (see a comprehensive review in Pironon *et al.*, 2017). This hypothesis assumes that the geographic central area is the site where conditions are more favourable and stable (Brown, 1984), and where species should be more abundant or have better performance, decreasing to the geographic periphery (Brown, 1984; Sexton *et al.*, 2009; Abeli *et al.*, 2020). A central assumption of CPH is that geographical and environmental spaces are concordant (Brown, 1984); however, in recent years, predictions derived from the CPH are not the commonly found patterns (Sagarin and Gaines, 2002; Abeli *et al.*, 2014; Santini *et al.*, 2019). The main explanation for the absence of center-periphery patterns under CPH predictions is the widely accepted statement that the geographical and environmental spaces are not concordant (Martínez-Meyer *et al.*, 2013; Van Couwenberghe *et al.*, 2013; Santini *et al.*, 2019), *i.e.*, not all ecologically marginal populations are located on the geographical periphery and vice versa (Soulé, 1973). More recent studies have found that population dynamics is associated with distance to the ecological centre, also called niche centroid (*e.g.* Martínez-Meyer *et al.*, 2013; Weber *et al.*, 2017; Osorio-Olvera *et al.*, 2019).

Species performance has been extensively studied along a geographical and ecological centre-periphery gradient, focusing on the current distribution of the focal species (*e.g.* Carey *et al.*, 1995; Nantel and Gagnon, 1999; Jump and Woodward, 2003; Costa *et al.*, 2016), and on the current ecological environment, more recently based on Ecological Niche Modelling (ENM) (*e.g.* Martínez-Meyer *et al.*, 2013; Weber *et al.*, 2017; Osorio-Olvera *et al.*, 2019). However, these studies do not take into account historical dimensions of species range, such as past environmental conditions (*e.g.* from the Last Glacial Maximum; LGM). These conditions, which change through time, create stable/unstable areas that could affect species traits that respond to long-term environmental conditions, consequently influencing species

performance. Therefore, there is a need to incorporate a historical approach to species performance studies (Keppel *et al.*, 2012; Tzedakis *et al.*, 2013; Pironon *et al.*, 2015). Further, ENM is a tool to identify climatically stable areas for long periods of time, under a historical niche perspective. Indeed, reconstruction of palaeo- and current distributions through ENM, together with the study of genetic patterns is a multidisciplinary framework to identify stable areas (*e.g.* Waltari *et al.*, 2007; Collevatti *et al.*, 2012; Baranzelli *et al.*, 2017).

The historical approach has been recently used in genetic (*e.g.* Eckert *et al.*, 2008; Duncan *et al.*, 2015; Pironon *et al.*, 2015) and demographic (*e.g.* abundance, germination and growth rate; Pironon *et al.*, 2015; Douda *et al.*, 2019) studies, evidencing its importance in significantly explaining species performance patterns (*e.g.* Pironon *et al.*, 2015; Douda *et al.*, 2019). Nonetheless, other important traits, such as morphological ones, remain unexplored from this perspective. Historically stable areas preserve the evolutionary history of lineages (Keppel *et al.*, 2012; Tzedakis *et al.*, 2013), and could harbor populations (and communities) where morphological traits are expressed differently from unstable areas (*e.g.* Myking and Yakovlev, 2006; Hatziskakis *et al.*, 2011; Varsamis *et al.*, 2020), including some traits related to the species performance. Variation or changes of some morphological traits may be linked to past species distribution rather than current ecological conditions (Douda *et al.*, 2019), especially traits that have long-term response to environmental conditions, or are strongly influenced by abiotic factors modeled over long periods of time (such as edaphic factors that influence tree height).

The study of morphological traits that determine species' performance has fundamental implications, both in the field of evolutionary ecology and in conservation biology (Sagarin *et al.*, 2006). Traits that influence performance are important resources to find high genetic diversity or geographic locations of genetic uniqueness (Lessica and Allendorf, 1995; Lira-Noriega and Manthey, 2014; Macdonald *et al.*, 2017), to describe the relationships between

species and the community (Fulton, 1999; Clark *et al.*, 2011; Lebrija-Trejos *et al.*, 2016), to understand environmental preferences (Díaz *et al.*, 2004; Wright *et al.*, 2004; Moles *et al.*, 2009; Marks *et al.*, 2016; Baranzelli *et al.*, 2020), and also as descriptors of the environment of a population or species (Díaz and Cabido, 2001), among other aspects. Specifically, functional traits in plants, such as leaf size, longevity, and canopy height and structure, are subject to environmental selection pressures (Díaz and Cabido, 2001); therefore, they are a fundamental determinant of species performance, in general related to environmental stability and productivity (Violle *et al.*, 2007), that should be explored under the historical centreperiphery approach. Unlike other traits that influence species performance, such as demographic, genetic or physiological traits, morphological functional traits are useful to understand environmental selective pressures and climatic changes more directly (Díaz and Cabido, 2001; Violle *et al.*, 2007), and to compare attributes (*i.e.* measurements) of each trait from distant ecosystems (Reich *et al.*, 1997; Díaz *et al.*, 2004); additionally, obtaining those attributes arises as an easy and inexpensive method (Pérez-Harguindeguy *et al.*, 2013).

In this study, we propose the inclusion of morphological traits in centre-periphery studies adding a historical approach to the geographical and ecological ones. Specifically, we present the case of the forest tree *Bulnesia sarmientoi* Lorentz ex Griseb. (Zygophyllaceae), an endangered species, distributed in central-northern Argentina, southeast Bolivia, western Paraguay, and marginally in Brazil (Zuloaga *et al.*, 2008; Waller *et al.*, 2012). We selected *B. sarmientoi* as a model species because a previous phylogeographical and niche modelling study (Camps *et al.* 2018) evidenced that genetic diversity, past demography and distribution range of the species were significantly affected by Pleistocene climatic fluctuations. Moreover, a Pleistocene refugium for *B. sarmientoi* was proposed in a climatically stable area from the LGM up to the present, which harbours the highest values of genetic diversity (Camps *et al.*, 2018). Additionally, a previous ecological study revealed a significant

variation in diameter, height and population density of *B. sarmientoi*, suggesting differences in growth dynamics across its range (Loto *et al.*, 2018).

Based on the previous phylogeographical study of *B. sarmientoi* (Camps *et al.*, 2018) showing a significant imprint of past climatic changes on the evolutionary history of the focal species, our general hypothesis is that the historical processes would have shaped variations in *B. sarmientoi* tree and leaf traits; therefore, the historical approach is expected to be a better predictor of population performance than the approaches that consider the geographical or ecological space, finding differences in morphological traits between the historical center, representing a climatically stable area, and the historical periphery. To test our hypothesis, we used and compared three approaches (geographical, ecological, and historical) to define central and peripheral populations, and examined which of them best explains morphological variation in *B. sarmientoi*. The ecological (*i.e.* geographical distance to the current niche centroid) and the historical (*i.e.* geographical distance to the LGM niche centre) approaches were based on Camps *et al.* (2018).

MATERIALS AND METHODS

Study area and species data

Bulnesia sarmientoi is a large tree species (up to 20 m) (Fig. 1); it is dominant in heavy clayrich soils, with highly impeded drainage and temporary anaerobiosis (Adamoli *et al.*, 1972; Prado, 1993). The species is distributed in central-northern Argentina, southeast Bolivia, western Paraguay, and marginally in Brazil (Zuloaga *et al.*, 2008; Waller *et al.*, 2012) (Fig. 2). The distribution area mainly includes the Dry Chaco ecoregion and marginally the Humid Chaco, Chiquitano Dry Forest, and Pantanal, according to Olson *et al.* (2001) classification. Temperature in the *B. sarmientoi* distribution range can reach an annual mean close to 26°C, whereas the minimum annual mean temperature in the area is 22°C [MERRAclim database]

(Vega *et al.*, 2017)]; the entire region is a warm, seasonal environment. In contrast, the precipitation gradient is more marked, with higher rainfall in the north (1200 mm) and southeast (850 mm), and lower in the centre-west (450 mm) [Worldclim database (Hijmans *et al.*, 2005)] (Fig. 2). The precipitation gradient ranges from semi-arid to humid, determining different ecoregions and ecotone zones (Cabrera, 1976; Olson *et al.*, 2001).

Sampling and trait measurements

The morphological traits of stem, canopy (both hereafter called "tree traits") and leaves (hereafter called "leaf traits") of mature individuals of Bulnesia sarmientoi were studied in 24 populations. Sampled populations covered the entire distribution range of *B. sarmientoi*, except for the Pantanal ecoregion populations. At this point it is important to mention that the few samples from the northern portion of the range reflect the natural distribution and abundance of the species, so it is not a sampling bias. The collecting procedure and minimum number of samples was performed following Pérez-Harguindeguy et al. (2013). To obtain tree traits, digital photographs of the five tallest mature individuals per population (118 trees total; in two populations only four trees were measured) were taken in a straight line to the tree, with the lens pointing towards the trunk, at breast height. Next to the trunk, a 150 cm wooden plank (demarcated every 50 cm) was placed (see Fig. 1B), which served as a reference to measure the tree traits in the software HOJA 3.6 (Verga, 2015). The tree traits measured were: canopy length (LC), canopy width (WC), total height (ToH), trunk height (TrH), trunk diameter (D), total height/canopy width ratio (RHC) and total height/trunk diameter ratio (RHD) (Fig. 2). These traits were measured because B. sarmientoi showed variation in size (e.g. diameter and height) according to the type of forest where they grew (Loto et al, 2018). In addition, height is recognized as a trait indicating resource availability (Fulton, 1999; Marks et al., 2016), while canopy structure showed intraspecific variation associated to environmental conditions and interspecific competition (*e.g.* Benavides *et al.*, 2019; Lang *et al.*, 2019; Ding *et al.*, 2020).

Five leaves per tree, from five trees per population, were collected and measured (600 leaves in total). The leaves were selected randomly from the middle portion of different branches that were in intermediate parts of the treetop, whenever possible. Digital images obtained from scanned leaves were used to take leaf measurements using HOJA 3.6. The leaf traits measured were: dry mass (DM), leaf area (LA), leaf length, leaf width (LW), petiole length (PL), specific leaf area (SLA), total leaf area (TLA), length/width ratio (RLW) (Fig. 2). Measured leaf traits were selected based on previous studies suggesting SLA variation is associated with resource availability (Díaz *et al.*, 2004; Wright *et al.*, 2004), and leaf size variation with moisture availability or aridity (*e.g.* Kleinschmit, 1993; Souza *et al.*, 2018; Li *et al.*, 2020). To obtain DM, the leaves were oven-dried at 70°C for 80 hours. SLA was calculated as a one-sided area of a fresh leaf, divided by its oven-dry mass.

Geographical approach

The geographical approach consisted of evaluating how tree and leaf traits vary across the distribution range with respect to the geographical centre (GC) and the geographical periphery (GP). The GC was calculated with the polygon centroid tool, implemented with SAGA 2.3.2 in Quantum GIS 2.18 (QGIS Development Team, http://www.qgis.org/). The centre has an area of 61,930 km², which is located equidistant to the north/south and east/west largest dimensions of the total distribution range of *B. sarmientoi* (412,864 km²). Each population was classified as GC or GP (Fig. 3).

Ecological approach

The ecological approach consisted of evaluating how tree and leaf traits vary in two components related to the niche centroid, calculated from current climatic conditions. Two methods were used to calculate the niche centroid: the first one was calculated from Ecological Niche Modeling (ENM) analysis, and the other from Mahalanobis and Euclidean distances. We consider the calculation of two different niche centroid because ENM are widely used to test CPH, but the distances seem to be better indicators of environmental suitability, mainly the Mahalanobis distance (Osorio-Olvera *et al.*, 2019).

The niche centroid calculation derived from an ENM analysis of *B. sarmientoi* previously reported (Camps *et al.*, 2018). In that study, seven algorithms were tested and the best three were selected according to the validation metrics (Camps *et al.*, 2018). We used those three best algorithms (Table 1) to obtain a niche centre of the species considering a probabilistic range. The niche centre was calculated as a range that represented the highest 15% probability of each algorithm (Table 1), taking the threshold as the minimum value of that range, and the highest probabilistic value as the maximum value. The ranges (*i.e.* niche centre) inferred with the three algorithms were overlapped on a map to obtain a consensus area representing the climatically most favourable area for the species (Fig. 3). Then, each studied population was classified as niche centroid (NC_ENM) when it was included in the consensus area, or as niche periphery (NP_ENM) when it was outside the favourability area.

The second niche centroid was calculated using the Mahalanobis and Euclidean distances, also taken from Camps *et al.* (2018), and was based on the distance to the fundamental niche of the species, defined by the three variables that most contributed to the ENM mentioned above (mean temperature of most humid quarter, specific humidity mean of coldest quarter, and annual mean specific humidity). The average of Mahalanobis and Euclidean distances to the climatic centroid, called suitability (varying between 0 and 1), was used to classify the

studied populations as central or peripheral (Fig. 3). Populations with suitability values greater than 0.85 were considered to be in the niche centroid (NC_ME), and populations with suitability values less than 0.85 were considered to be in the niche periphery (NP_ME).

A buffer area of 6.28 km (maximum diagonal distance of one pixel) was considered for these two niche centroid calculations. Thus, any sampled population that was found outside the centroid but less than 6.28 km away from that area, was considered part of the niche centroid. The buffer distance was calculated in Quantum GIS 2.18.

Historical approach

The historical centre-periphery approach consisted of defining a historical central area (HC) based on previous phylogeographic and paleodistribution analyses (Camps *et al.*, 2018). In that study, a putative climatic refugium for *B. sarmientoi* was proposed; this area has been climatically stable from the LGM up to the present and is currently a hotspot of genetic diversity (Camps *et al.*, 2018). In the same way as the ecological approach, a buffer area of 6.28 km was considered. Thus, we considered populations to be in the historical centre (HC) if they were found in the area of the climatic refugium, or to be in the historical periphery (HP) if they were found outside the refugium (Fig. 3).

Statistical analyses

Measured tree and leaf traits were standardized and only those traits not strongly correlated (Pearson correlation value <0.8) were included for subsequent analyses (see Supplementary data Figs S1 and S2). To assess the presence of distance-based patterns of variation in morphological traits, a spatial autocorrelation analysis was performed for each leaf and tree trait. Significance levels of Moran's I coefficients of spatial autocorrelation were obtained using Monte Carlo methods with 999 simulations. Three different distances were

tested, according to the minimum, average, and maximum distance that separate the studied populations (40 km, 70 km and 100 km, respectively). These analyses were performed in InfoStat v.2018 (Di Rienzo *et al.*, 2018).

For each centre-periphery approach, a Principal Component Analysis (PCA) and a Linear Mixed-Effects Models (LMM) were performed, each with one set of morphological traits (*i.e.* tree and leaf). PCA was used to obtain new, fully uncorrelated variables, which synthesize the information from the original data set (McGarigal *et al.*, 2000). For the PCA, the *princomp* methodology and the *ggbiplot* library (Vu, 2016) were used in R 3.4.4 (R Core Team, 2018) through RStudio 1.1.4 (RStudio Team, 2015). The number of components to be retained was based on Parallel Analysis (Horn, 1965) with 5000 iterations (adjusted eigenvalue > 1) using the *paran* R package (Dinno, 2018). The retained components of the PCA were saved and used in the LMM.

A LMM was performed to test which of the implemented centre-periphery approaches significantly explained morphological patterns, using the new variables obtained with the PCA as response variables, and the population location (centre vs. periphery) as a fixed factor. A mixed effects model was chosen to incorporate and control the absence of independence of the data from the same population, using 'population' as a random factor. LMM was performed using the *nlme* library (Pinheiro *et al.*, 2020) of R 3.4.4 through RStudio 1.1.4. Each retained component of leaf and tree variation presented normal distribution (Shapiro-Wilks test P > 0.05; the residues were adjusted approximately to the theoretical normal line).

RESULTS

Morphological traits analyses

The tree traits measured in *Bulnesia sarmientoi* that were not strongly correlated (Pearson correlation value < 0.8) were: tree height/canopy width ratio (RHC), height/trunk diameter ratio (RHD), total height (ToH), and trunk height (TrH). On the other hand, the leaves traits that did not show a strong correlation (Pearson correlation value < 0.8) were: dry mass (DM), leaf length (LL), petiole length (PL), length/width ratio (RLW) and specific leaf area (SLA). Correlation matrices and a summary of the morphological measures obtained for each sampled population are shown in Supplementary data Figs S1 and S2 and Table S1, respectively). The spatial autocorrelation analysis indicated that none of the measured morphological traits (*i.e.* tree and leaf traits) show a significant distance-based pattern (Moran's I p value > 0.05), in any of the three tested spatial distances.

For the PCA performed to obtain the new synthetic variables from tree traits, two components were retained, representing 80.9% of the total tree trait variation. ToH and TrH traits made the greatest contribution to principal component (PC) 1 (-0.656 and -0.605, respectively), and the RHC trait to PC 2 (0.807). For the PCA of leaf traits, two PCs were retained, summarizing 68% of the total variance. DM and LL were the most variable traits along PC 1 (-0.594 and -0.565, respectively) and SLA was the most variable along PC 2 (0.825).

Geographic centre-periphery

Climatically, the geographical centre was a dry warm environment, while the periphery was a warm rainy environment (see Fig. 2). Geographical central and peripheral populations showed a differentiation in the multivariate space of tree traits (Fig. 4A). The biplot of the PCA shows trees from central localities to the right of the plots, presenting lower values of

TrH and ToH than those located on the periphery. However, the LMM showed no significant differences between the centre and periphery in tree traits, on either PC axis (Table 2). Regarding leaf variation, the PCA presented a differentiation between the geographical centre and the periphery in the multivariate space (Fig. 5A). The biplot of the PCA shows those individuals that belong to the geographical centre mainly differentiated from peripheral populations in relation to PC 2, indicating small SLA values in the geographical centre. LMM confirmed the significant differences between centre and periphery in the leaf traits (PC 2) (Table 2, Fig. 6B).

Ecological centre-periphery

The centre derived from the ENM was mainly dry, and contains the lowest temperatures in the species range (see Fig. 2). The periphery derived from the ENM was warmer and wetter towards the north and east. Conversely, the centre derived from the Mahalanobis and Euclidean distances showed a north-south gradient of temperature and precipitation, being the north warmer and drier than the south (see Fig. 2). Meanwhile, the periphery presented temperature and precipitation values representative of the range of values found across the species' distribution area, without specific climatic conditions. Regarding the PCA performed with tree traits, ecological central and peripheral populations were not differentiated in the PCA biplot (Fig. 4B, C). LMMs confirmed that there were no differences between populations located in the ecological centre compared to those located in the same way, the PCA performed with leaf traits, did not show a clear differentiation in traits between central and peripheral populations (Fig. 5B, C). LMMs confirmed that there were no differentiation in traits between

Historical centre-periphery

Besides being a climatically stable area this historical centre presented, both during the LGM and current times, lower precipitation levels than the historical periphery (see Supplementary data Fig. S3, Table S2 and Fig. S4). For the PCA performed with tree traits, historical central and peripheral populations showed a differentiation in the multivariate space (Fig. 4D). The biplot of the PCA showed trees from central localities to the right of the plots, presenting lower values of TrH and ToH than those located on the periphery. LMMs confirmed the significant differences between centre and periphery in the PC1 of the tree traits (Table 2, Fig. 6A). Conversely, the PCA performed with leaf traits, did not show a clear differentiation in traits between central and peripheral populations (Fig. 5D). LMM confirmed no significant differences between the centre and periphery in leaf traits, on either PC axis (Table 2).

DISCUSSION

In this study, we investigated which of three centre-periphery approaches better explains the morphological differentiation of the tree species *Bulnesia sarmientoi*. The spatial localization of the three centres considered were not coincident, as was also reported in previous studies (*e.g.* Pironon *et. al.*, 2015). In particular, the non-correspondence between the geographic location of ecological and historical centres, suggests that the location of the niche optimum has changed over time. The historical centroid approach evidenced a centre-periphery pattern only for tree traits, showing smaller trees in the historical centre than in the historical periphery. Otherwise, we found significant differences in leaf traits between the geographical

centre and the periphery, mainly due to smaller specific leaf area values toward the geographical centre with respect to the periphery. Previous studies had already reported that a historical approach significantly explained genetic and demographic traits (*e.g.* Pironon *et al.*, 2015; Douda *et al.*, 2019), but no previous studies have used these approaches to analyse morphological traits determinant of species performance.

Historical and geographic approaches explain traits variation: possible processes underlying the retrieved centre-periphery patterns

Spatially, the geographic, ecological and historical centre-periphery approaches were not concordant. Given these spatial and climatic differences, not all approaches explained traits variation. Climatically, there were similarities across approaches; the geographic, ecological and historical centres were drier than peripheral areas, however they did not coincide in the temperature ranges, being the geographic centre the warmest of them. The tree trait pattern obtained using the historical approach showed lower values of total height and trunk height towards the historical centre than towards the historical periphery. The historical centre encloses a putative climatic refugium for B. sarmientoi, which is currently a hotspot of genetic diversity and has been a climate-stable area from the Last Glacial Maximum (LGM) up to the present (Camps et al., 2018). Regarding environmental conditions, LGM refugium (*i.e.* historical centre) persists in sites where the soil was formed by Andean sediments from the Pliocene (Ramos and Ghiglione, 2008; Iriondo, 2010), and where the driest climate conditions in the entire range have occurred over time (see Supplementary data Fig. S3, Table S2 and Fig. S4). Interestingly, according to field observations, most sites within the proposed LGM refugium (i.e. historical centre) showed a special type of forest commonly named "palosantales"; this association with palosantales was not identified with the other approaches. In palosantales, which are usually monospecific and with a higher population density than in mixed forests (see Fig. 1A, B), *B. sarmientoi* individuals showed much smaller trunk diameter and did not develop the size observed in mixed forests (Loto *et al.*, 2018). Thus, the historical component explaining tree trait variation would be related to the persistence of this kind of forest in the historical centre. This could be either due to two not mutually exclusive factors that could have been modulated by historical processes: the lack of competition due to the absence of other forest species in this stressful area, and the direct effect of stressful conditions affecting tree traits. In this line, previous studies showed tree height is lower in low precipitation sites (Fulton, 1999; Moles *et al.*, 2009). In addition, low height may indicate low soil fertility and richness (Fulton, 1999; Marks *et al.*, 2016). Ongoing studies of species that coexist with *B. sarmientoi* in mixed forests, such as *Aspidosperma quebracho-blanco* and *Prosopis* spp., will enhance the understanding if this refugium was a favourable site only for *B. sarmientoi*.

According to the three centre-periphery approaches studied here, the geographical centre approach is the one that best represents the precipitation gradient, from a dry warm environment in the geographical centre to a warm rainy environment towards the periphery. The centre derived from the ecological approach was dry, but its temperature was highly variable, having areas that presented the highest temperature values and others that presented the lowest temperatures values across the species range; therefore, the ecological centre as a whole, did not represent an "extreme" environment. The ecological approach (and not by the ecological approach) the obtained pattern showing the highest specific leaf area values in the geographical periphery makes sense, since it is indicative of high resource availability and productive environments towards peripheral populations (Reich *et al.*, 1998; Albert *et al.*, 2010). Low specific leaf area values indicate dense leaves and low growth rates, and are associated with resource conservation and dominance in areas of stressful conditions (Díaz *et*

al., 2004; Wright *et al.*, 2004; Albert *et al.*, 2010). This environmental response of specific leaf area is current and not historical (*i.e.* it was not revealed using the historical approach), which is consistent with the expected phenotypic plasticity of functional traits (see Díaz and Cabido, 2001; Wright *et al.*, 2004; Violle *et al.*, 2007).

The environmental characteristics of the geographic and historical centre-periphery approaches (i.e. low precipitation, high temperatures and resource-poor soils), and the observed trait patterns, show that B. sarmientoi is a stress-tolerant species. Furthermore, the mean value of specific leaf area for the studied populations was low to intermediate (see Supplementary data Table S1) compared to specific leaf area measurements taken at global scale (Reich et al., 1998; Pérez-Harguindeguy et al., 2013), suggesting that it is a conservative species (sensu Díaz et al., 2004); that is, it would have a conservative strategy in the use of resources, with low performance and productivity rates. On the other hand, in resource-rich environments, such as those occupied by the species in the Humid Chaco, Chiquitano Dry Forest or Pantanal (all peripheral environments under the geographic and historical approaches), tree height or the specific leaf area present values associated with a better performance (high tree height and high specific leaf area). However, our field observations registered low population abundance in these peripheral sites, suggesting that interspecific competition could be conditioning species performance. It seems that in the geographic and historical peripheral sites individual traits are maximized, while in central sites (stressful environments) population performance is maximized.

The differential pattern between morphological traits and species abundance highlights that the former are better indicators of environmental pressures than species abundance, thus, there is a need to include them more frequently in centre-periphery studies (Dallas *et al.*, 2017; Santini *et al.*, 2019). Although we analyzed the tree and leaf traits as two data sets, and not as individual traits, we detected that the statistical differences were strongly influenced

mainly by the tree height and the specific leaf area, as mentioned above. These results are consistent with previous studies suggesting both traits have a great response to environmental conditions (Reich *et al.*, 1998; Fulton, 1999; Díaz *et al.*, 2004; Moles *et al.*, 2009; Albert *et al.*, 2010; Cosacov *et al.*, 2014). Instead, the remaining measured traits which did not show a strong centre-peripheral pattern, probably are less variable at intraspecific level (Siefert *et al.*, 2015) or they could be variable at a different spatial scale from the one considered in this study, associated with other environmental factors (*e.g.* soil properties, or environmental variables not considered here).

On the absence of some centre-periphery patterns

In disagreement with other studies (*e.g.*, in population abundance, Martínez-Meyer *et al.*, 2013; Van Couwenberghe *et al.*, 2013; Weber *et al.*, 2017) our results show that the ecological approach is a poor predictor of centre-periphery patterns, despite having tested two niche centroids derived from Ecological Niche Modelling output and from Mahalanobis and Euclidean distances. However, we know that an ecological approach does not necessarily explain traits variation, as reviewed for population abundance (Santini *et al.*, 2019). Probably, as suggested by Dallas *et al.* (2017), there are important factors other than the environment in the regulation of population dynamics, such as dispersal boundaries and unmeasured ecological interactions. It should be taken into account that the Ecological Niche Modelling (ENM), used to define the niche centroid, is based on climatic similarities rather than on fitness (Osorio-Olvera *et al.*, 2019). Another important consideration is that ENM is a current "snapshot" of a particular moment in the history of the focal species, and the environment it occupies today may not be the one that maximizes its fitness. These theoretical limitations of the ecological centre-periphery approach are also applicable to the historical approach, since the latter is methodologically derived from the former. However, a key difference should be

considered: areas that are climatically stable over time are informative about the persistence of the species in a specific geographical area, and therefore the approach is useful in demographic terms (see evidence of climatically stable areas inferred by ENM and genetic demographic analysis in Carstens and Richards, 2007; Waltari *et al.*, 2007; Collevatti *et al.*, 2012; Baranzelli *et al.*, 2017, among others). Climatic stability (*i.e.* the historic centre) resulted in an area where the environmental conditions persisted over time and left their imprint, especially in tree traits that have long-term environmental response. Hence, we suggest that the historical approach performs better than the ecological one in reflecting the suitability of a geographical area for a species. Future research should explore other methodologies to define centres and peripheries under an ecological approach, considering key factors shaping the form and function of morphological traits.

Another outstanding result was the lack of significant differences in tree traits in the geographical centre with respect to the geographical periphery. In the geographical central area, the most extreme climatic conditions or low resources occur, being these conditions reported as causes of low values for total height (Marks *et al.*, 2016). One possible explanation is the time scale, because tree height variation was associated with long-term processes, in opposition to more plastic traits, like leaf size or specific leaf area (traits that evidenced a geographic centre-periphery pattern). On the other hand, the absence of a geographic gradient in tree traits is possibly the consequence of the fact that this gradient does not detect the optimal microsites for the species. These microsites present environmental characteristics, different from the surrounding, which positively affect the performance of a species (Dunwiddie and Martin, 2016; Mayoral *et al.*, 2019). We consider that, unlike the other approaches, the historical approach returns these microsites because the historical center is inferred based on long term climatic suitability (with genetic patterns corroborating the long lasting persistence of the species in that area), and climate is the main structuring

factor (Keppel *et al.*, 2012) affecting also other environmental factors such as edaphic conditions. But also, the absence of geographical gradient in tree traits may be due to the advanced process of deforestation and selective logging in the Gran Chaco (Mereles and Pérez de Molas, 2008; Hansen *et al.*, 2013; Vallejos *et al.*, 2015; Pometti *et al.*, 2021). These threats, which are particularly intense in the mixed forests, could reduce the chances of sampling high trees.

CONCLUSIONS

Our results show that each trait data set (*i.e.* tree and leaf trait) studied in *Bulnesia sarmientoi* showed a different pattern, explained by different approaches. The geographic approach explained leaf trait variation, probably associated with environmental suitability; the geographic centre presents the most severe climatic conditions of the distribution area, since it is an area characterized by dry conditions and high temperatures. The historical approach allows the recovery of historical processes underlying tree traits variation. In the historical centre a greater number of palosantales forests are present; historically this center has been the driest area across the species geographic range. These patterns observed for the two morphological data sets suggest that both sets respond to environmental processes following a centre-periphery dynamic; while tree trait variation was associated with long-term processes, leaf trait patterns were associated with short-term conditions. The ecological approach was a poor predictor of centre-periphery patterns, probably because the measured traits could not capture the effect of meaningful ecological processes. Due to the dynamic nature of the geographical ranges, delimitation of the centre and the periphery should be based not only on current geographical and ecological perspectives, but also on historical ones.

Based on our research, we highlight the use of a historical perspective in the explanation of centre-peripheral patterns, and the inclusion of morphological traits to understand species performance across their range, in addition to demographic, physiological or genetic traits. Our results showed that the historical approach could elucidate historical processes underlying morphological traits variation, reinforcing the importance of this approach and the use of morphological functional traits to understand environmental selective pressures. Future studies could explore centre-periphery morphological patterns under the three approaches used in our study in co-distributed species, to elucidate if there is a geographical, ecological ts

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FIGURE CAPTIONS

Fig. 1. *Bulnesia sarmientoi* in its natural habitat. A) "palosantal", i.e. a particular vegetation type usually monospecific and conformed by small trees. B) Single or very few tall *B. sarmientoi* trees in a degraded mixed forest. C) Immature fruit. D) Leaves.

Fig. 2. Distribution of *Bulnesia sarmientoi*, bioclimatic variables, and morphological traits measured across the species distribution range. A) Annual mean temperature. B) Annual precipitation. C) Traits measured on trees. D: trunk diameter; LC: canopy length; ToH: total height; TrH: trunk height; WC: canopy width. D) Traits measured on leaf. LL: leaf length; LW: leaf width; PL: petiole length. Bioclimatic variables obtained from MERRAclim (Vega *et al.*, 2017) and Worldclim database (Hijmans *et al.*, 2005). Population codes are given in Supplementary data Table S1.

Fig. 3. Maps of centre-periphery approaches explored in *Bulnesia sarmientoi*. A) Geographical approach, B) Ecological approach based on Ecological Niche Modelling analysis, C) Ecological approach based on Mahalanobis and Euclidean distances to climatic centroid, and D) Historical approach. *B. sarmientoi* distribution was delimited by Camps *et al.* (2018). Population codes are given in Supplementary data Table S1.

Fig. 4. Principal Component Analysis graphics of tree traits. A) Geographical approach, B) Ecological approach based on Ecological Niche Modelling analysis, C) Ecological approach based on Mahalanobis and Euclidean distances to the climatic centroid, and D) Historical approach. The contribution of each trait is proportional to the length of its corresponding vector. RHC: height/canopy width ratio; RHD: height/trunk diameter ratio; ToH: total height; TrH: trunk height. GC: geographical centre; GP: geographic periphery; NC_ENM: niche

centroid; NP_ENM: niche periphery; NC_ME: niche centroid; NP_ME: niche periphery; HC: historic centre; HP: historical periphery.

Fig. 5. Principal Component Analysis graphics of tree traits. A) Geographic approach, B) Ecological approach based on Ecological Niche Modelling analysis, C) Ecological approach based on Mahalanobis and Euclidean distances to climatic centroid, and D) Historical approach. The contribution of each trait is proportional to the length of its corresponding vector. DM: dry mass; LL: leaf length; PL: petiole length; RLW: length/width ratio; SLA: specific leaf area. GC: geographical centre; GP: geographical periphery; NC_ENM: niche centroid; NP_ENM: niche periphery; NC_ME: niche centroid; NP_ME: niche periphery; HC: historical centre; HP: historical periphery.

Fig. 6. Linear Mixed-Effects Models graphics. Only the LMMs that indicated significant differences (P < 0.05) are shown. A) LMM of tree trait PC1 in historical climatic centre-periphery approach, and B) LMM of leaf trait PC2 in geographical centre- periphery approach. GC: geographical centre; GP: geographical periphery; HC: historical centre; HP: historical periphery.

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TABLES

Table 1. Minimum value (threshold), greater probabilistic value, and 15% upper range of the three algorithms used for the *Bulnesia sarmientoi* Ecological Niche Modelling.

Algorithm	Threshold	Greater probabilistic value	15% upper range		
Bioclim (Busby, 1991)	0.013	0.759	0.647 to 0.759		
Support Vector Machine (Vapnik, 1998)	0.027	0.040	0.038 to 0.040		
Maxent (Phillips et al., 2006)	0.115	0.717	0.627 to 0.717		
cer's	60				

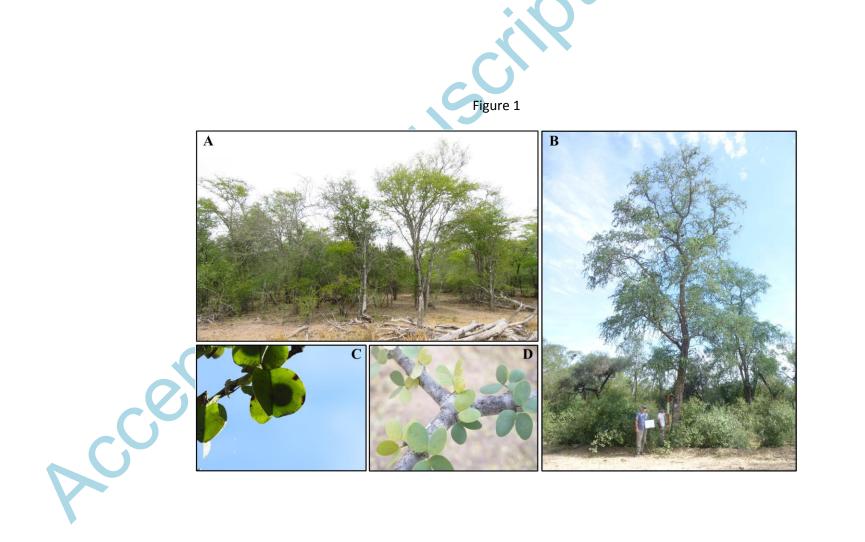
Table 2. Results of the Linear Mixed-Effects Models to test the effect of centre-periphery in

 leaf and tree traits summarized on Principal Component axes.

Approach and variables	Tree traits				Leaf traits							
	Estimate	SE	df	<i>t</i> value	<i>p</i> value	Estimate	SE	df	<i>t</i> value	<i>p</i> value		
1. Geographical												
PC1	0.66	0.52	22	1.3	0.22	0.13	0.52	22	0.25	0.80		
PC2	0.115	0.33	22	0.35	0.73	0.76	0.36	22	2.1	0.048		
2a. Ecological (Ecological Niche Modelling)												
PC1	0.7	0.38	22	1.9	0.077	0.56	0.37	22	1.5	0.15		
PC2	0.8	0.24	22	1.6	0.12	0.058	0.29	22	0.2	0.84		
2b. Ecological (Mahalanobis and Euclidean distances)												
PC1	0.36	0.42	22	0.84	0.41	-0.41	0.40	22	-1.01	0.32		
PC2	-0.089	0.26	22	-034	0.74	0.48	0.29	22	1.6	0.12		
3. Historical												
PC1	1.06	0.34	22	3.1	0.005	0.161	0.39	22	0.41	0.68		
PC2	0.37	0.24	22	1.6	0.13	0.32	0.29	22	1.12	0.27		

Note: Significant effects ($P \leq 0.05$) are shown in bold. PC1: principal component 1; PC2:

principal component 2. SE: standard error; df: degrees of freedom.



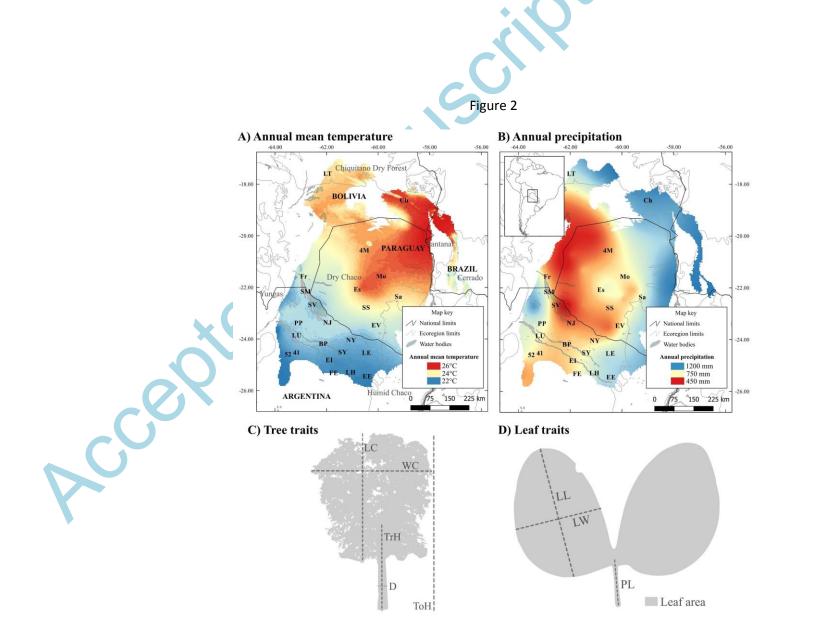
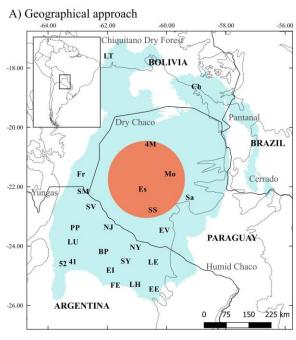
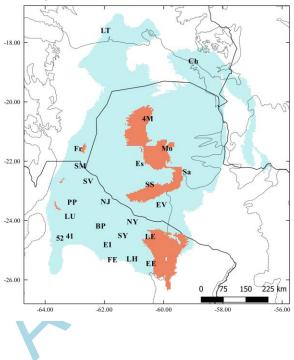
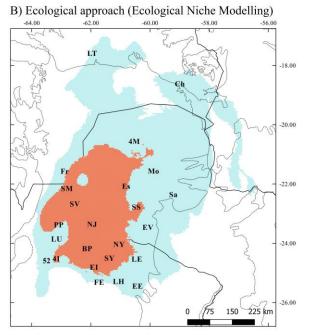


Figure 3

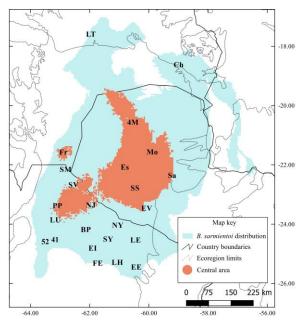


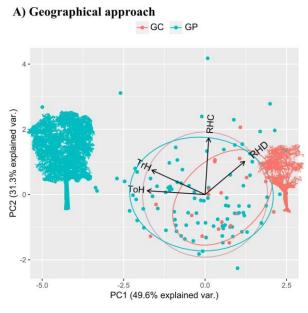
C) Ecological approach (Mahalanobis and Euclidean distance)





D) Historical approach



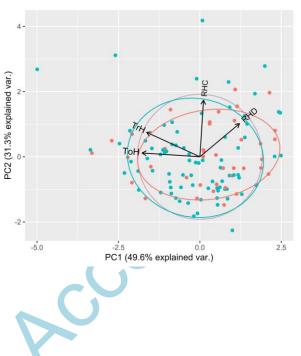


4-(1) 2--2--5.0 -2-5 0.0 -2.5 PC1 (49.6% explained var.)

B) Ecological approach (Ecological Niche Modelling)

- NC_ENM - NP_ENM

C) Ecological approach (Mahalanobis and Euclidean distance)



D) Historical approach

