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Geographic variation in the morphology of the sand-dwelling lizard *Nothobachia ablephara* (Squamata: Gymnophthalmidae)

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Abstract

Geographic variation in the morphology of the sand-dwelling lizard Nothobachia ablephara (Squamata: Gymnophthalmidae). Nothobachia ablephara is a small microteiid lizard with an elongated body and reduced limbs; it occurs in isolated dune fields in the state of Bahia (Xique-Xique and Alagoado) and small sandy patches in northeastern Brazil. A previous molecular study found a marked mtDNA divergence between populations of N. ablephara from Alagoado and Xique-Xique dunes, suggesting that the two populations diverged from one another between 3 and 4 million years ago. Given this isolation, it is interesting to explore whether morphological traits of the lizards reflect the reported genetic divergence of the populations. Scale counts of the sexes and the populations differ significantly, but there is considerable overlap of values. Univariate and multivariate analyses revealed significant morphometric variation between sexes and populations; however, this is mostly explained by size differences. Females are larger than males in all characters that are sexually dimorphic, and individuals from Xique-Xique are larger than those from Alagoado in all characters that vary geographically. The sample from Alagoado has more sexually dimorphic characters than the one from Xique-Xique. Although N. ablephara displays some geographical variation, the two populations could not be unequivocally distinguished by scale counts and morphometric data.

Keywords: caatinga, differentiation, dunes, morphometry.

Resumo

Variação geográfica na morfologia do lagarto fossorial *Nothobachia ablephara* (Squamata: Gymnophthalmidae). *Nothobachia ablephara* é um pequeno microteídeo de corpo alongado e membros reduzidos que ocorre em campos de dunas isolados no estado da Bahia (Xique-Xique e Alagoado) e em pequenas manchas de areia na região nordeste do Brasil. Dados moleculares mitocondriais encontraram uma divergência significativa entre as populações de *N. ablephara* dos campos de dunas de Xique-Xique e Alagoado, sugerindo que essas duas populações divergiram uma da outra entre três e quatro milhões de anos atrás. Considerando que as populações estão isoladas, é

interessante explorar se caracteres morfológicos desses lagartos refletem a divergência entre as duas populações reportada pelos dados moleculares. As contagens de escamas mostraram diferenças significantes entre sexos e entre as populações, mas há uma significativa sobreposição dos valores. Análises univariadas e multivariadas revelaram uma variação significativa na morfometria entre sexos e populações que foi principalmente explicada por diferenças de tamanho. Fêmeas são maiores do que os machos em todos os caracteres que são sexualmente dimórficos, e os indivíduos de Xique-Xique são maiores do que os de Alagoado em todos os caracteres que mostraram diferenças geográficas. Ademais, a amostra de Alagoado apresentou dimorfismo sexual em maior número de caracteres que a de Xique-Xique. Embora *N. ablephara* tenha exibido variação geográfica significativa, não foi possível distinguir inequivocamente as duas populações com base em contagens de escama e morfometria.

Palavras-chave: caatinga, diferenciação, dunas, morfometria.

Introduction

Microteiid lizards of the family Gymnophthalmidae (Estes et al. 1988) comprise about 250 species in 48 genera of small to mediumsized lizards (snout-vent length ranging from 3-15 cm) that range from southern Mexico to Argentina (Uetz and Hosek 2018). Most microteiids are terrestrial to semi-fossorial and occupy a wide range of habitats, from open areas in the Andes to lowland rain forests (Pianka and Vitt 2003). Many species are microhabitat specialists—e.g., semi-aquatic (Gelanesaurus, Neusticurus, and Potamites), scansorial (Anadia, Placosoma, and Selvasaura) or fossorial taxa (Bachia, Calyptommatus, Nothobachia, and Scriptosaura) (Rodrigues 1984, 1991a, Ávila-Pires 1995, Rodrigues et al. 2001, Rodrigues and Santos 2008, Moravec et al. 2018); thus, there is convergence of body forms in distantly related groups (Pellegrino et al. 2001, Wiens et al. 2006, Marques-Souza et al. 2018).

The occurrence of convergent characters, especially the recurrent appearance of morphological adaptations to fossoriality, combined with the paucity of many taxa in collections impose difficulties for a morphology-based taxonomy of Gymnophthalmidae (Rodrigues 1991a, Pellegrino *et al.* 2001). The first contribution to the molecular systematics of Gymnophthalmidae was based on data for 51

species representing 26 genera (Pellegrino *et al.* 2001). Subsequently, new molecular studies have provided a deeper understanding of the relationships between species and genera of Gymnophthalmidae (Rodrigues *et al.* 2001, Castoe *et al.* 2004, Doan and Castoe 2005, Goicoechea *et al.* 2016, Marques-Souza *et al.* 2018).

Taxonomy is an important component of biogeography, ecology, conservation, and other biodiversity studies (Ebach et al. 2011). Historically, herpetological studies focused on morphological data to assess the relationships among taxa, and traditional characters such as cephalic scales, number of dorsal and ventral scales, presence or absence of morphological structures and color pattern were commonplace for species recognition in microteiids (Rodrigues 1984, 1991a-c, Rodrigues and Santos 2008, Recoder et al. 2014). In the last few decades the fields of systematics and taxonomy have been revitalized and transformed by the confluence of phylogenetic thinking and more universal access to the tools of molecular analysis (Padial et al. 2010).

Nothobachia is monotypic, containing only *N. ablephara* Rodrigues, 1984. The species is characterized by an extremely elongated body with reduced limbs, absence of eyelids and external ear opening, and fossorial and diurnal habits. Within the Gymnophthalmini tribe, the

morphological and molecular data indicate that *N. ablephara* is closely related to two other genera, *Calyptommatus* and *Scriptosaura*, which in addition to other characters, share striking reduction of limbs, absence of eyelids and an external ear opening (Rodrigues 1991a, Pellegrino *et al.* 2001, Rodrigues and Santos 2008, Goicoechea *et al.* 2016). *Nothobachia ablephara* is endemic to semi-arid Caatinga of northeastern Brazil and occurs in the sand dune fields of Xique-Xique and Alagoado and in small sandy patches in southern Piauí and Pernambuco states, west of the San Francisco River.

Siedchlag et al. (2010) sought to understand phylogeographic patterns in fossorial microteiid lizards in the dunes of São Francisco based on molecular data of the mitochondrial cytochrome b gene (cyt b) and segments of the ribosomal genes 12S and 16S. These authors found a marked divergence (i.e., 6-8% in combined mtDNA) between populations of Nothobachia ablephara from the dune fields of Xique-Xique and Alagoado. This divergence was about ten times greater than the divergence found among samples from three localities within the Xique-Xique dune field. This difference in mtDNA suggests that the northern and southern populations of N. ablephara, which are separated by over 100 km of unsuitable habitat, diverged between three and four million years before the present (Siedchlag et al. 2010). This same pattern of differentiation was observed between populations of Eurolophosaurus divaricatus (Rodrigues, 1986) (Tropiduridae), which also is restricted to dune fields on the left bank of São Francisco river (Passoni et al. 2008). Molecular data for one specimen of N. ablephara from sandy patches in Petrolina, Pernambuco state, was used to place the species in the Gymnophthalmidae phylogeny (Pellegrino et al. 2001, Goicoechea et al. 2016), but these data were not used to test divergence among populations (Siedchlag et al. 2010).

Given the high molecular divergence found between populations of *Nothobachia ablephara* and the importance of a well-resolved taxonomy for biodiversity studies, we undertake a detailed morphological study of specimens from different populations of this species to test whether morphological differences exist that reflect the genetic divergence, and whether the differences are sufficient to warrant the recognition of distinct species.

Materials and Methods

Study Area

The sand dunes are located along the middle São Francisco River Valley in the Brazilian morphoclimatic domain of semiarid Caatinga and represent a paleodesert in northeastern Brazil (Rodrigues 1996). Mean annual temperatures exceed 26.2°C, the highest of which is in the state of Bahia, and mean annual rain fall ranges from 400-800 mm; typically, the dry season extends from April-September (Rocha and Rodrigues 2005). The dunes are variable in elevation (20-100 m a.s.l.) and the soils are composed mainly of eolic sand (Rodrigues 1991a, 1996). The vegetation on the dunes is greenish year-round, even in the dry season, and differs from that on the neighboring Caatinga where the appearance of the vegetation changes completely during the dry season (Rodrigues 1991a).

Lizards were collected from each of the two major sand-dune systems on the São Francisco River (Figure 1); there are molecular data documenting the divergence between the two populations. Alagoado (09°29' S, 41°21' W) is in the north and Xique-Xique (10°33' S, 43°19' W) in the south; the localities are separated by a region of rocky/clayish soils extending for approximately 150 km (Rodrigues 1991a).

Data Collection

We sampled a total of 112 adult *Nothobachia ablephara* from the two major sand dune fields in this study (Ramiro *et al.* 2017)—34 from the south (Xique-Xique) and 78 from the north

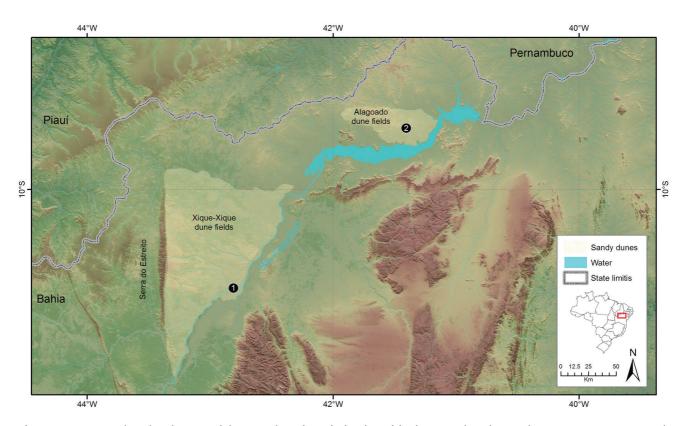


Figure 1. Geographic distribution of the samples of *Nothobachia ablephara* used in this study: (1) Xique-Xique (south population) and (2) Alagoado (north population), Bahia state.

(Alagoado). In addition, we examined all of the individuals housed at the herpetological collections of Museu de Zoologia da Universidade de São Paulo (MZUSP) and Laboratório de Herpetologia, Instituto de Biociências da Universidade de São Paulo (MTR), São Paulo (Appendix I).

Six measurements (to nearest 0.1 mm) were taken with digital calipers from left side of preserved specimens: snout-vent length (SVL), from posterior border of cloaca to the tip of snout; trunk length (TL), between the anterior margin of hind limb to the posterior margin of forelimb; head height (HH), at highest point in the longitudinal axis; head width (HW), at the greatest distance between the temporal scales; head length (HL), from the junction of infra and supralabial scales to the tip of snout and hind-limb length (HLL), from the insertion of the limb to the tip of the longest finger. Four scales counts were also recorded: number of scales

around mid-body (SAB); number of pairs of gular scales (GUL); and numbers of transverse rows of dorsals (DOR) and ventrals (VEN). Male lizards were identified by the presence of preanal pores in males (Rodrigues 1984), and the total number of pores (POR) was recorded.

Data Analysis

We log₁₀ transformed the measurements prior to analysis to meet requirements of normality (Bailey and Byrnes 1990, Zar 2010). The log-transformed morphometric data were tested for normality with Shapiro Wilk's Test and equivalence of variances with Levene's Test (Zar 2010). We assessed the Pearson correlation between the morphometric variables for both sexes in both localities.

The scale counts DOR, VEN, and SAB, deviated from normality (Shapiro Wilk's test, p < 0.05). Thus, to evaluate differences in scale

counts between sexes and between populations, we used the non-parametric Mann-Whitney U Test. We compared the number of gular scales (GUL) and the number of preanal pores (POR) qualitatively because little variation was observed. The arrangement of gular scales varied between even rows of gulars (symmetric) or uneven rows of gulars (asymmetric); frequencies in each population were recorded.

Because differences in sample sizes can affect statistical results and we had fewer specimens from the south (Xique-Xique), we follow a methodology suggested by Cardini et al. (2015) to evaluate the impact of sample size bias in our significance tests using the SVL of specimens from the Alagoado Dune Field. First, we checked for the presence of outliers in the samples of males and females using boxplots (not shown). For females (N = 42) and males (N = 42)= 36), the mean and the variance were calculated for the complete sample and for 10 random samples with N = 5, N = 10, N = 15, N = 20, N= 25, and N = 30. The 61 mean and variance values calculated were plotted for each category and each sex to visualize differences between values, and then we conducted a Mann-Whitney U Test to evaluate whether statistical differences in significance existed between the differentsized samples.

All morphometric variables have a strong linear correlation (R > 0.6) except the length of the hind limb (HLL) (R < 0.6). Differences in hind-limb length (HLL) were tested with twoway analysis of variance (ANOVA) using sex and population as grouping factors. Pairwise comparisons in HLL were made using a pooled one-way ANOVA (4 groups) and post-hoc significance accessed via Tukey's HSD tests. The remaining morphometric variables, with R coefficients higher than 0.6, were used in a principal component analysis (PCA) summarize the variability in morphometric data in order to remove among-variables correlation. The PC scores that together accounted for most variation (> 95%) were saved and used to highlight differences between both populations

and sexes in those morphometric variables, with a multivariate analysis of variance (MANOVA) using the PC scores as dependent variables and sex and population as grouping factors.

We performed all statistical analyses using IBM SPSS Statistics (version 20.0, SPSS Inc., 2011).

Results

The comparative morphometric data for females and males of both populations of Nothobachia ablephara are presented in Table 1. There is sexual dimorphism in DOR and VEN (Mann-Whitney U Test, U = 0.014) in both populations, and females have more scales than males in both localities (Table 1). When comparing the two populations, females differed only in VEN (Mann-Whitney U Test, U = 0.018) and males differed in VEN (Mann-Whitney U Test, U = 0.001) and DOR (Mann-Whitney U Test, U = 0.001) scales. Females and males from the southern population have more scales than do males and females of the northern population. There were no sexual or geographic differences in SAB (Mann-Whitney U Test, U < 0.16).

There usually are seven or eight gular scales (GUL) in symmetrical rows. The northern population has seven or eight pairs of gulars in both sexes and asymmetries varying from 6-9 scales on each side. The frequency of asymmetries in this population is 0.27 for females and 0.21 for males. Males of the southern population have seven or eight pairs, but all females have seven pairs. There were asymmetries in both sexes, varying from 6-8 scales on each side. The frequency of asymmetries in this population is 0.41 for females and 0.5 for males. Males of both populations typically have six preanal pores (POR) (N = 32 north, N = 13 south), but some individuals have five pores (N = 3 north, N = 2)south) or four pores (N = 1 north, N = 2 south).

In our evaluation of impact of sample size bias in significance tests, we found no significant differences between the two statistical parameters

Table 1. External morphological measurements (mm) and scale counts (average ± standard deviation) of females and males of two populations of *Nothobachia ablephara*. *N* = sample size, HL = head length, HH = head height, HW = head width, TL = trunk length, SVL = snout-vent length, HLL = hind limb length, DOR = number of dorsal scales, VEN = number of ventral scales and SAB = scales around the body.

	Nothobachia ablephara (north)		Nothobachia ablephara (south)	
	Females $(N = 42)$	Males $(N = 36)$	Females ($N = 17$)	Males $(N = 17)$
HL	4.9 ± 0.4	4.5 ± 0.3	4.9 ± 0.6	4.7 ± 0.3
HH	2.3 ± 0.2	2.2 ± 0.2	2.3 ± 0.3	2.2 ± 0.2
HW	3 ± 0.3	2.7 ± 0.2	2.9 ± 0.2	2.9 ± 0.2
TL	40.5 ± 3.9	36.1 ± 3.8	39.8 ± 4.7	35.8 ± 8.3
SVL	53.3 ± 4.9	47.6 ± 3.9	53.1 ± 6.1	48.9 ± 3.3
HLL	5.7 ± 0.5	5.3 ± 0.4	6.1 ± 0.5	5.9 ± 0.6
DOR	45 ± 1.2	42.7 ± 0.8	45.3 ± 0.9	44.3 ± 1.2
VEN	37.3 ± 0.9	35.3 ± 0.8	37.9 ± 1.3	36.5 ± 0.9
SAB	19.6 ± 0.9	19.3 ± 0.9	20.1 ± 0.9	19.7 ± 0.6

(mean and variance) estimated for SVL, among samples with different sample sizes (N > 5) for either females (mean: Mann-Whitney U Test, p = 0.971; variance: Mann-Whitney U Test, p = 0.063) or males (mean: Mann-Whitney U Test, p = 0.156; variance: Mann-Whitney U Test, p = 0.358) (Figure 2). Thus, the samples were pooled in two geographic groups for subsequent analysis—Alagoado (N = 42 females, N = 36 males) and Xique-Xique (N = 17 females, N = 17 males).

Hind-limb length (HLL) varies significantly between sexes (ANOVA, F = 9.44, p = 0.003) and populations (F = 22.07, p < 0.001), but sex and population did not interact significantly (F = 0.30, p < 0.05). In pairwise comparisons, females from the northern population have longer hind limbs than males (Tukey's HSD test, p < 0.01), but sexes from southern population did not differ in HLL (p = 0.433) (Figure 3). Both females and males have significant geographic differences (p < 0.05), and lizards from the southern population have longer hind limbs than do the northern lizards (Figure 3).

The first principal component explains 80.85% of the variation in all individuals, with high and positive loadings for all morphometric

variables (HL, HH, HW, LBL and SVL) (Table 2). The second component accounts for 7.53% of the variation with negative loadings for head length (HL), length between limbs (LBL) and snout–vent length (SVL). The third component explains only 6.43% of the variation, and the fourth, 4%. In the morphological space of the PCA (Figure 4) the first component subtly separates females and males independent of the population, with most of the females having lower scores than males, but neither of the two components separates sexes or populations clearly.

The multivariate analysis using the first four principal components differ significantly between populations (MANOVA, p = 0.005), between sexes (MANOVA, p = 0.0001), and there is an interaction between sex and geography (MANOVA, p = 0.004).

Discussion

Our results indicate that *Nothobachia* ablephara has significant geographical variation in morphology. Nevertheless, there is a general overlap in the distribution of values of the quantitative characters from Xique-Xique and

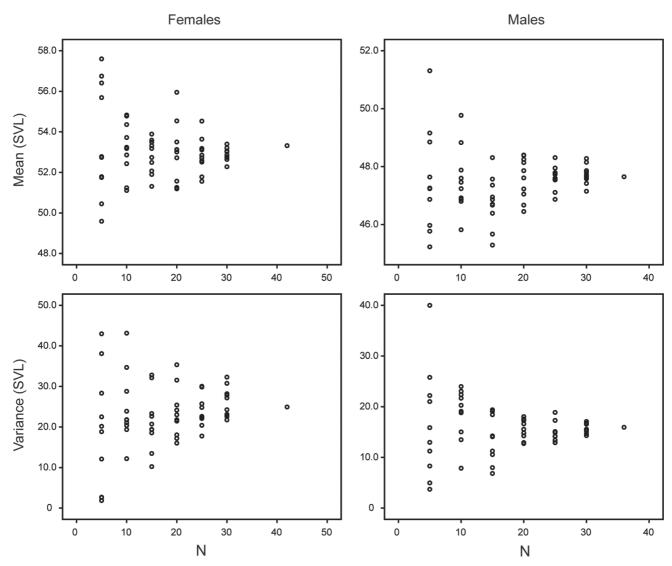


Figure 2. Effects of sample size on statistical parameters. Estimates refer to mean (upper) and variance (lower) of SVL measured for females and males of north population, based on 10 replicates for each of six subsample categories (N = 5, 10, 15, 20, 25, 30) and total sample (N = 42 and 36, respectively). See text for details.

Alagoado. Therefore, the quantitative data we used in this study, which have proven useful for the taxonomy of many genera of gymnophthalmini lizards (Rodrigues 1991a–c, Ávila-Pires 1995, Recoder *et al.* 2014, Rodrigues *et al.* 2017), provide no diagnostic characters that unambiguously distinguish the two populations. Despite the significant divergence in the mDNA between the populations, we are not proposing taxonomic alterations in *Nothobachia* and consider it a single species (pending additional

data) because the populations cannot be distinguished by diagnostic morphological characters.

Based on different studies of geographically separated organisms having a similar pattern of high molecular divergence and absence of morphological diagnostic characters, we hypothesize three major possible explanations: (1) despite long-term isolation of populations, speciation has not occurred because similar ecological conditions and low selective pressures

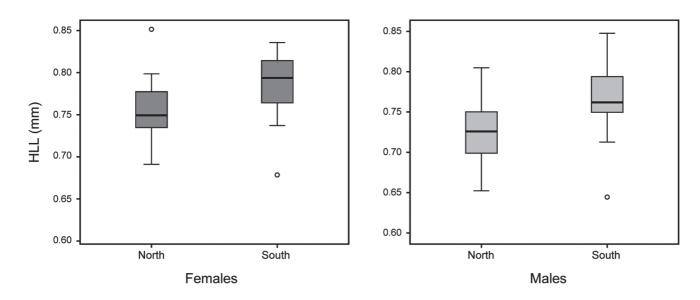


Figure 3. Boxplot of hind-limb length measurements (HLL) for females (dark gray) and males (light gray) from north and south populations.

prevail; (2) cryptic speciation has occurred; or (3) owing to secondary contact between individuals from both populations, gene-flow occurred.

Environmental conditions might impose stabilizing selection on morphology (Bickford *et al.* 2006); both populations of *Nothobachia ablephara* are under similar climatic conditions in sand dunes. Both Alagoado and Xique-Xique are characterized by high mean annual

temperatures and unpredictable rain regimes, even during the wet season. However, natural history data are only available for the southern population from Xique-Xique sand dunes (Rocha and Rodrigues 2005). Thus, more ecological traits need to be studied to determine whether environmental conditions and/or niche variation have had a role in the maintenance of external morphology in populations of *N. ablephara*. Based on the possibility that environmental

Table 2. Loadings of the first four principal components of a PCA and the percentage of variation of each component for populations of *Nothobachia ablephara*. HL = head length, HH = head height, HW = head width, TL = trunk length and SVL = snout–vent length.

	PC1	PC2	PC3	PC4
HL	0.855	-0.254	0.445	0.080
HH	0.871	0.408	-0.012	0.274
HW	0.899	0.257	0.084	-0.344
TL	0.921	-0.225	-0.274	0.042
SVL	0.947	-0.170	-0.203	-0.040
% Variation	80.85	7.53	6.43	4.06
Cumulative %	80.85	88.39	94.82	98.87

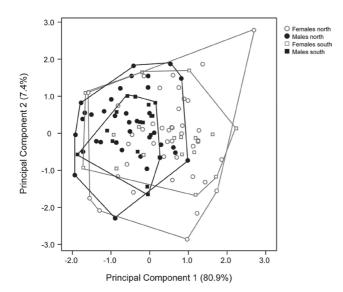


Figure 4. Results of principal component analysis (PCA) on six morphometric measurements of two samples of *Nothobachia ablephara*. Circles correspond to samples from the northern population, and squares to the southern population. Empty symbols correspond to females, filled symbols correspond to males.

conditions might impose stabilizing selection on morphology, it is important to notice that the other two closely related genera of fossorial lizards, Calyptommatus and Scriptosaura, also conserved morphologies. recognized species of *Calyptommatus*, all living in sand patches around the middle São Francisco River, have highly similar external morphologies and are diagnosed mainly by head scalation and differences body-scale subtle in texture. Scriptosaura catimbau Rodrigues and Santos, 2008 is the sister taxon of N. ablephara and despite their different generic allocations, the species are similar in size, body shape, and color pattern.

During the past three decades, molecular data have revealed the existence of complexes of cryptic species for which it has been difficult to identify diagnosable morphological characters (Bickford *et al.* 2006, Siler *et al.* 2011), a pattern that recently seemed to be common in tropical lizards (Nunes *et al.* 2012, Domingos *et al.* 2014,

Werneck et al. 2015, Guarnizo et al. 2016, Teixeira Jr. et al. 2016). The identification of cryptic species relies largely on genetic data (Bickford et al. 2006); thus, it is important to test whether the potential cryptic lineages have restricted hybridization and complete isolation 2000). and Jockusch Frequently, mitochondrial data have been used to determinate cryptic lineages in other lizard species (Harris et al. 2002, Pinho et al. 2007, Nunes et al. 2012, Domingos et al. 2014). The divergence in mitochondrial data in Nothobachia ablephara (> 6%) is similar to that observed for morphologically distinct species of Calyptommatus (> 3.6%) and lower than the divergence between populations of C. leiolepis Rodrigues, 1991 from Alagoado and Xique-Xique (< 4.6%) (Siedchlag et al. 2010). Thus, cryptic diversification may also be a potential explanation for the pattern observed in N. ablephara, with high divergence in mitochondrial data but subtle morphological

In some cases, hemipenial morphology is a key taxonomic character in investigations of cryptic speciation because of it has higher rates of evolution than do other morphological traits (Klaczko et al. 2015, D'Angiolella et al. 2016). In other species having hemipenial variation among different populations, further molecular data revealed cryptic species complexes coincident with the morphology of the hemipenis (Inger and Marx 1962, Zaher and Prudente In *Iphisa* elegans Grav. (Gymnophthalmidae), analysis of the morphology of the hemipenis revealed five distinct morphotypes that correspond to five wellsupported molecular clades (Nunes et al. 2012). Gabirot et al. (2010, 2012, 2013) investigated the differences in chemical sexual signs between populations of *Podarcis hispanicus* (Steindachner, 1870) (Lacertidae) and their importance in the process of reproductive isolation and cryptic speciation. In P. hispanicus, males of five populations secreted a singular and characteristic mixture of compounds used as sexual signals that can classify them. Data about the anatomy of the hemipenis of *Nothobachia ablephara* are only available for specimens from Alagoado (Nunes *et al.* 2014). Despite the lack of data about mating strategies, *N. ablephara* is a fossorial species, does not show sexual dimorphism in color pattern, and only males present preanal pores, that produce a large amount of secretion, indicating that the chemical communication could be an important aspect of *N. ablephara* life history. It would be interesting to test if the hemipenial morphology or the chemical characterization of the preanal glands secretions vary interpopulationally.

Given the current absence of nuclear molecular data for N. ablephara, we cannot determine whether the populations experienced gene-flow more recently than the time of divergence of 3-4 m.y. suggested by the mitochondrial genes (Siedchlag et al. 2010). Geological evidence suggests that the sand dune fields were formed largely in the Pleistocene during a drier, more arid climate with intense eolian activity (Tricart 1974), and that dunes are inactive and fixed in their current positions by vegetation for at least 28,000 years (Barreto 1996, Oliveira et al. 1999). Thus, we could assume that the Xique-Xique and Alagoado populations of Nothobachia have been isolated at least since the end of the Pleistocene. given the small body size of these lizards and the fidelity of the species to sand soils (Rodrigues 1984), it seems unlikely this species could disperse across about 150 km of typical, hardsoiled caatinga that currently separates the dune fields. Nuclear data should be analyzed to exclude the hypothesis of gene-flow postmitochondrial divergence.

In conclusion, interpopulational morphological differentiation in external characters is not significant *Nothobachia ablephara*, and there are insufficient molecular data to address the hypotheses on the population divergence and evolutionary history of this species. The need for an integrative, multi-trait approach that brings information about chemical composition, hemipenial morphology, fine-scale anatomy,

resource use, and reproductive behavior is obvious and critical to an understanding the factors involved in geographical differentiation in *N. ablephara* and other morphologically conserved gymnophthalmid lizards.

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Appendix I. Specimens examined.

Nothobachia ablephara: Brazil. Bahia. Alagoado (MZUSP: 71469, 71486, 71487, 71490–71495, 71499, 71516–71521, 71523–71525, 71566, 71529–71535, 71543, 71567, 71568, 71570, 71572, 74885, 75991, 75992, 75994, 75995, 75998–76000, 76003, 76004, 76007–76010, 77870, 77871, 77948–77951, 77953–77957, and 91915–91921; MTR: 21291, 21292, and 21294–21304). Brejo da Umburana (MZUSP: 80217–80220, and 80222). Ibiraba (MZUSP: 71471, 71473–71477, 71480, and 95114). Queimadas (MZUSP: 76869–76874, 76876, 76878, 77872–77875, 77940, and 77943–77947). Barra (MTR: 39038 and 39039).