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ANOLIS CHRYSOLEPIS DUMÉRIL AND BIBRON, 1837 (SQUAMATA: IGUANIDAE), REVISITED: MOLECULAR PHYLOGENY AND TAXONOMY OF THE *ANOLIS CHRYSOLEPIS* SPECIES GROUP

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ABSTRACT. The *Anolis chrysolepis* species group is distributed across the entire Amazon basin and currently consists of *A. bombiceps* and five subspecies of *A. chrysolepis*. These lizards are characterized by moderate size, relatively narrow digital pads, and a small dewlap that does not reach the axilla. We used the mitochondrial gene *ND2* to estimate phylogenetic relationships among putative subspecies of *A. chrysolepis* and taxa previously hypothesized to be their close relatives. We also assessed the congruence between molecular and morphological datasets to evaluate the taxonomic status of group members. On the basis of the two datasets, we present a new taxonomy, elevating each putative subspecies of *A. chrysolepis* to species status. We provide new morphological diagnoses and new distributional data for each species.

Key words: *Anolis*, Amazon, Iguanidae, molecular phylogeny, taxonomy

RESUMO. O grupo de espécies *Anolis chrysolepis* atualmente consiste em *A. bombiceps* e cinco subespécies de *A. chrysolepis*, ocupando toda a Bacia Amazônica. Esses lagartos são caracterizados por tamanho moderado, lamelas digitais relativamente estreitas e um papo extensível que não chega às axilas. Nós utilizamos o gene mitocondrial *ND2* para estimar as relações filogenéticas entre as subespécies de *A. chrysolepis* e táxons previamente considerados parentes próximos. Nós também determinamos a congruência entre conjuntos de dados morfológicos e moleculares, para avaliar o status taxonômico dos membros desse grupo. Com base nos dois conjuntos de dados, apresentamos uma nova taxonomia, elevando

cada subespécie de *A. chrysolepis* ao status de espécie. Fornecemos novas diagnoses morfológicas e novos dados de distribuição para cada espécie.

Palavras-chave: *Anolis*, Amazônia, Iguanidae, filogenia molecular, taxonomia

INTRODUCTION

The Pleistocene Refuge Hypothesis proposed almost simultaneously by Haffer (1969) and Vanzolini and Williams (1970) posits that patches of lowland tropical forest that existed during dry periods in the Pleistocene served as core areas for speciation in birds and in the lizard complex *Anolis chrysolepis*, respectively. Although the Pleistocene Refuge Hypothesis has been falsified for members of the *A. chrysolepis* species group because diversification occurred much earlier (15 mya) than the Pleistocene (Glor et al., 2001), relationships among all members of the group have not been worked out and related taxa (e.g., *A. meridionalis* and *A. bombiceps*) have not been properly placed with reference to the *A. chrysolepis* complex, and current names do not accurately reflect the evolutionary history of the group (Glor et al., 2001; Nicholson et al., 2005). Because the *A. chrysolepis* species group has been and continues to be a model for evolutionary (Nicholson et al., 2006, 2007; Schaad and Poe, 2010) and ecological (Vitt and Zani, 1996; Vitt et al., 2001, 2008) studies, it is critical that their relationships be properly understood. Here we present a phylogenetic hypothesis for the *A. chrysolepis* species

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group using a much larger set of samples than was available previously and provide species names for taxa that can be identified as independent evolutionary lineages. Following de Queiroz (2007), we consider independent evolutionary lineages, here recognized on the basis of gene trees, analogous to species. Results of this study should be directly applicable to phylogeographic and phyloecological studies of the *A. chrysolepis* species group.

The *A. chrysolepis* group comprises two species: *A. chrysolepis* Duméril and Bibron, 1837, and *A. bombiceps* Cope, 1876. *Anolis chrysolepis* is currently composed of five subspecies: *A. chrysolepis chrysolepis*, in eastern Guiana (Brazil, French Guiana, Suriname, and southern Guyana); *A. chrysolepis planiceps* Troschel, 1848, in western Guiana (Brazil, Suriname, northwestern Guyana, Venezuela, and Trinidad); *A. chrysolepis scypheus* Cope, 1864, in western Amazonia (Colombia, Ecuador, Peru, and northwestern Brazil); *A. chrysolepis tandai* Avila-Pires, 1995, in southwestern Amazonia (Brazil and Peru); and *A. chrysolepis brasiliensis* Vanzolini and Williams, 1970, in Brazil, from Maranhão and enclaves of open vegetation in southern Pará south to São Paulo (Vanzolini and Williams, 1970; Avila-Pires, 1995; Icochea et al., 2001; Santos-Jr et al., 2007). *Anolis bombiceps* occurs in western Amazonia, in Peru, Colombia, and Brazil, at least in partial sympatry with *A. c. scypheus* and perhaps also with *A. c. tandai* (Avila-Pires, 1995). Members of the *A. chrysolepis* group are characterized by their moderate size (up to 83 mm snout–vent length); short heads; supraorbital semicircles usually forming a pronounced ridge; relatively narrow digital pads, with distal lamellae under phalanx ii forming a slightly prominent border; a dewlap that does not reach the axilla and is present in both sexes (but smaller in females); and keeled, imbricate ventral scales that are distinctly larger than dorsals.

The *A. chrysolepis* species group was examined morphologically by Vanzolini and Williams (1970), who recognized four sub-

species of *A. chrysolepis* and a distinct species, *A. bombiceps*. Vanzolini and Williams (1970: 13) believed the level of differentiation between the subspecies were “closest to species difference, and indicative, perhaps, of past and future potential species formation.” *Anolis chrysolepis* was later examined by Avila-Pires (1995) under the name *A. nitens*. She described another subspecies, *A. n. tandai*, and observed that most specimens occurring in areas of intergradation according to Vanzolini and Williams (1970) could be assigned to one of the recognized subspecies.

Very little subsequent taxonomic research has been conducted on the species of the *A. chrysolepis* group. One molecular phylogenetic study included three of the described *A. chrysolepis* subspecies and found they formed a weakly supported clade (Glor et al., 2001). Glor et al. (2001: 2664) concluded that, “further study of geographical genetic interactions among these subspecies probably will reveal that they are distinct species.” Additional molecular phylogenetic research, with broad outgroup sampling, recovered a well-supported clade consisting of *A. onca*, *A. annectans*, *A. lineatus*, *A. auratus*, *A. meridionalis*, and *A. chrysolepis*, although *A. chrysolepis* was represented by just a single individual from Roraima, Brazil (Nicholson et al., 2005). Members of this clade were included in another phylogenetic analysis (Nicholson et al., 2006), using the same three *A. chrysolepis* subspecies of Glor et al. (2001), which recovered a paraphyletic *A. chrysolepis*. Nicholson et al. (2006) found that *A. c. tandai* was more closely related to *A. meridionalis* and the *A. onca* + *A. annectans* clade, whereas *A. c. scypheus* and *A. c. planiceps* formed a clade that was the sister group to the remaining species + *A. auratus*. Like Glor et al. (2001), Nicholson et al. (2006) stressed the need for additional research into the systematics of *A. chrysolepis* and the possible existence of cryptic species. *Anolis bombiceps* has not been included in any molecular studies so far.

The name *A. chrysolepis* has a long and confusing history, with both *A. nitens* and *A.*

chrysolepis considered valid names for the species (Hoogmoed, 1973; Avila-Pires, 1995; Myers and Donnelly, 2008). Myers and Donnelly (2008) presented a detailed history of the use of these names, and Myers (2008) requested the International Commission of Zoological Nomenclature (ICZN) to give precedence of *A. chrysolepis* Dumeril and Bibron, 1837, over *Draconura nitens* Wagler, 1830, which was accepted (ICZN, 2010).

In the present work, we analyzed mitochondrial DNA from the protein coding gene *ND2* and associated tRNA and morphological data from all five described subspecies of *A. chrysolepis* and related taxa to 1) recover the phylogenetic relationships among subspecies of *A. chrysolepis* and test previous phylogenetic hypotheses, 2) evaluate the taxonomic status of described subspecies of *A. chrysolepis*, and 3) present a revised taxonomy that incorporates this phylogenetic information.

MATERIALS AND METHODS

Taxon Sampling and DNA Sequencing

We sampled representatives of each of the five subspecies of *A. chrysolepis* (Table 1, Figure 1). Species previously shown to be closely related to *A. chrysolepis* were also included either from newly sequenced samples (e.g., *A. bombiceps*) or from previously published GenBank material (Glor et al., 2001; Nicholson, 2002; Nicholson et al., 2005, 2006). Genomic DNA was extracted from muscle, liver, or tail clips using DNeasy Blood and Tissue Kit (Qiagen, Valencia, California). Polymerase chain reaction was used to amplify portions of the mitochondrial protein-coding gene *ND2* (NADH dehydrogenase subunit 2) and adjacent tRNAs with primers LVT_Met-f.6_AnCr (AAGCTATTGGGCCCATACC) and LVT_5617_AnCr (AAAGTGYYTGTGATTCA) (Rodriguez Robles et al., 2007). Polymerase chain reaction cleanup and DNA sequencing was performed by Agencourt Bioscience (Beverly, Massachusetts). Sequences were edited and aligned

using SEQUENCHER ver. 4.2 (Gene Codes, Ann Arbor, Michigan). *ND2* sequences were translated into amino acids using MacClade ver. 4.08 (Maddison and Maddison, 1992) to confirm alignment and gap placement and ensure there were no premature stop codons.

Phylogenetic Analyses

We analyzed the *ND2* data using parsimony in PAUP ver. 4.0b10 (Swofford, 2001). Parsimony analysis was conducted using a heuristic search with 1,000 random taxon additions and tree bisection and reconnection (TBR) branch swapping and all characters equally weighted. We conducted 1,000 bootstrap replicates with 25 random additions per replicate to assess nodal support (Felsenstein, 1985).

Mitochondrial DNA (mtDNA) has been widely used to recover phylogenetic relationships among species and to delimit species (Avise et al., 1998; Grau et al., 2005; Gamble et al., 2008; Fenwick et al., 2009), and because of its shorter coalescent times, it is considered a good indicator of population history and species limits (Avise et al., 2000; Wiens and Hollingsworth, 2000; Wiens and Penkrot, 2002; Zink and Barrowclough, 2008; Barrowclough and Zink, 2009). However, the high substitution rate of mitochondrial DNA makes saturation, especially at third codon positions, a possible problem for accurate phylogenetic reconstruction (Jukes, 1987; Yoder et al., 1996; Glor et al., 2001; Hudson and Turelli, 2003). One way to minimize the effects of saturation is to use model-based phylogenetic methods like maximum likelihood (ML) and Bayesian analyses (Felsenstein, 1978; Jukes, 1987; Huelsenbeck et al., 2001; Lartillot et al., 2007). Additionally, the use of partitioned model-based analyses, with separate models of molecular evolution for each gene or codon, can minimize phylogenetic error (Bull et al., 1993; Lemmon and Moriarty, 2004; Nylander et al., 2004; Brandley et al., 2005). We conducted Bayesian analyses using MrBayes 3.1.2

TABLE 1. MATERIAL EXAMINED FOR THE MOLECULAR PHYLOGENETIC ANALYSIS, INCLUDING TAXON NAME, MUSEUM NUMBERS, SPECIMEN LOCALITY, AND GENBANK NUMBERS.

<i>Anolis</i> Taxon	ID No.	Locality	GenBank
<i>A. tandai</i>	MPEG 22285	Itaituba, Pará, Brazil	JN191547
	MPEG 25029	Juruti, Pará, Brazil	JN191546
	LSUMZ H-14098	Rio Ituxi, Amazonas, Brazil	JN191542
	MPEG 25060	Coari, Amazonas, Brazil	JN191545
	LSUMZ H-16398	Rio Solimões, Amazonas, Brazil	JN191543
	LSUMZ H-16474	Rio Solimões, Amazonas, Brazil	JN191544
	LSUMZ H-13599	Rio Juruá, Acre, Brazil	JN191548
<i>A. chrysolepis</i>	MPEG 26590	Trombetas, Pará, Brazil	JN191532
	MPEG 26568	Faro, Pará, Brazil	JN191530
	MPEG 26563	Acari, Pará, Brazil	JN191531
	BPN780	Ralleighvallen, Suriname	JN191534
	MPEG 26584	Maicuru, Pará, Brazil	JN191533
	BPN 1587	Saul, French Guiana	JN191536
	BPN 1979	Saul, French Guiana	JN191541
	BPN 1874	Nouragues, French Guiana	JN191540
<i>A. scypheus</i>	LSUMZ H-12592	Reserva Faunistica Cuyabeno, Sucumbios Province, Ecuador	AF337804
	LSUMZ H-12543	Reserva Faunistica Cuyabeno, Sucumbios Province, Ecuador	AF337802
	LSUMZ H-12989	Reserva Faunistica Cuyabeno, Sucumbios Province, Ecuador	JN191568
<i>A. planiceps</i>	LSUMZ H-12300	Rio Ajarani, Roraima, Brazil	JN227867
	BPN 1080	Kartabo, Guyana	JN191550
	BPN 1082	Kartabo, Guyana	JN191549
	BPN 228	Imbaimadai, Guyana	JN191551
	BPN 96	Kartabo, Guyana	JN191552
<i>A. brasiliensis</i>	CHUNB 45077	Caseara, Tocantins, Brazil	JN191558
	CHUNB 45075	Minaçu, Goiás, Brazil	JN191559
	CHUNB 08842	Parauapebas, Pará, Brazil	JN191557
	CHUNB 43282	Brasília, Distrito Federal, Brazil	JN191555
	CHUNB 34542	Novo Progresso, Pará, Brazil	JN191556
	CHUNB 27158	Mateiros, Tocantins, Brazil	JN227868
	CHUNB 11521	Palmas, Tocantins, Brazil	JN191565
	CHUNB 37528	São Domingos, Goiás, Brazil	JN191563
	CHUNB 37527	Paraná, Tocantins, Brazil	JN191562
	CHUNB 52471	Peixe, Tocantins, Brazil	JN191560
	GRC 16378	Alto Paraíso, Goiás, Brazil	JN191561
<i>A. auratus</i>	LSUMZ H-13928	Alter do Chão, Pará, Brasil.	JN191571
<i>A. bombiceps</i>	KU 222145	1.5 km N of Teniente Lopez, Loreto, Peru	JN191570
<i>A. fuscoauratus</i>	LSUMZ H-13566	Rio Juruá, Acre, Brazil	AF337786
<i>A. meridionalis</i>	LF166692	Reserva Mbaracayú, Canindeyu, Paraguay	AY909760
<i>A. lineatus</i>	LSUMZ H-5450	Netherlands Antilles	AF294287
<i>A. onca</i>	CIEZAH1156	Estado Falcón, Venezuela	DQ377357
<i>A. annectens</i>	CIEZAH1160	Estado Falcón, Venezuela	DQ377345
<i>A. sericeus</i>	LACM7069	Costa Rica	AY909778
<i>A. isthmicus</i>	MFO191	Mexico	AY909762
<i>A. laevis</i>	MVCFC12252	Guatemala	AY909756
<i>A. sagrei</i>	KdQ1797	La Habana, Cuba	AF337778
<i>A. utilensis</i>	LDW12480	Honduras	AY909785
<i>A. grahami</i>	JBL 250	Discovery Bay, Jamaica	AF055938
<i>A. loveridgei</i>	USNM10683	Honduras	AY909759
<i>A. uniformis</i>	n/a	Belize	AY909784
<i>A. crassulus</i>	MZFC6458	Mexico	AY909748
<i>A. carolinensis</i>	CCA 8051	Unknown	NC010972

(Huelsenbeck and Ronquist, 2001) on both the partitioned and unpartitioned datasets. Data were partitioned by codon with a fourth partition for tRNAs and the optimal partitioning strategy selected using Bayes Factors calculated from the harmonic mean likelihood values (Nylander et al., 2004; Brandley et al., 2005). We estimated the best fit model of sequence evolution for the data as a whole and for each partition separately using AIC scores in Modeltest (Posada, 2008). Bayesian analyses were initialized with a neighbor-joining tree and two separate analyses conducted for each partitioning strategy. Each analysis consisted of seven heated chains and one cold chain run for 2 million generations, with sampling every 1,000 generations. Post-burnin convergence was checked by visual inspection of likelihood values by generation using Tracer 1.5 (Rambaut and Drummond, 2009) and visual inspection of split frequencies using AWTY (Nylander et al., 2008). We also conducted partitioned Maximum Likelihood analysis, with data partitioned as above using RAXML ver. 7.0.4 (Stamatakis, 2006) using the GTR+GAMMA model for all partitions. We conducted 1,000 “fast bootstrap” replicates and 10 separate maximum likelihood searches. Bootstrap values ≥ 70 were considered as indicating strong support for both parsimony and ML analyses.

We calculated net among group distances (Nei and Li, 1979) between major lineages of the “*A. chrysolepis* species group” using MEGA 4 (Kumar et al., 2008). We calculated both uncorrected *p*-distances and corrected distances using the GTR model.

On the basis of our best ML tree, we compared alternative phylogenetic hypotheses using the Shimodaira-Hasegawa (SH) test (Shimodaira and Hasegawa, 1999) and the Approximately Unbiased (AU) test (Shimodaira, 2002). Three alternative hypotheses were considered: 1) monophyly of *A. chrysolepis* subspecies, excluding *A. bombiceps* and *A. meridionalis*; 2) monophyly of the *A. chrysolepis* subspecies + *A. bombiceps*, excluding only *A. meridionalis*;

and 3) monophyly of all *A. c. tandai* specimens, as identified by morphological data. We used RAXML7.0.4 (Stamatakis, 2006) to compute per-site log likelihoods that were input into CONSEL (Shimodaira and Hasegawa, 2001) to calculate *P* values. We also tested alternative phylogenetic hypotheses in a Bayesian framework and calculated the Posterior Probabilities of alternative hypotheses using the tree filter option in PAUP*.

Morphological Analyses

We collected morphological and morphometric data from 403 specimens (Appendix 1) from the following zoological collections: MZUSP, Museu de Zoologia da Universidade de São Paulo; CHUNB, Coleção Herpetológica da Universidade de Brasília; MPEG, Museu Paraense Emílio Goeldi; MCZ, Harvard Museum of Comparative Zoology; and KU, University of Kansas. Measurements were recorded with digital calipers to the nearest 0.1 mm on the right side of the body, except when specimens were damaged (in this case, the left side was used). Scale and measurement terminology follows Avila-Pires (1995).

We recorded the following morphometric data: snout-vent length (SVL), tail length (from posterior edge of preloacal plate), head width, head height, mouth length (from tip of snout to posterior margin of mouth), distance between orbits (minimum), ear-opening diameter, distance between nostrils (minimum), distance from mouth to ear (from anterior margin of ear-opening to posterior margin of mouth), snout length (from tip of snout to anterior margin of orbit), interparietal length, tibia length, foot length (from toe IV base to the heel), fourth toe length (from toe IV nail to toe base), and fourth toe maximum width. Additionally, we recorded the following meristic characters: scales around midbody, postrostrals, supralabials, infralabials, loreals (under second canthal), canthals, scales between second canthals, scales between supraorbital semicircles (minimum), scales

between interparietal and supraorbital semicircles (minimum), postmentals, fourth finger lamellae, and fourth toe lamellae.

A few measurements and scale counts could not be assessed for all specimens analyzed. In multivariate analysis, cases with missing observations will be dropped, weakening the analysis because of loss of information and degrees of freedom. To avoid simply deleting entire rows of data, missing observations can be estimated using a variety of methods, including mean substitution, regression, expectation maximization, maximum likelihood and multiple imputation (Tabachnick and Fidell, 2001; Quinn and Keough, 2002). Among these approaches for imputing values to missing observations, multiple imputation is the most robust and also makes fewer assumptions about the pattern of missing observations (Rubin, 1996; Van Buuren et al., 2006). Therefore, we imputed missing data using multivariate imputations by chained equations (Van Buuren et al., 2006), as implemented by package *mice* in R v. 2.12.0 (R Development Core Team, 2009).

To partition the total morphometric variation between size and shape variation, we defined body size as an isometric size variable (Rohlf and Bookstein, 1987) following Somers (1986): we calculated an isometric eigenvector, defined a priori with values equal to $p^{-0.5}$, where p is the number of variables (Jolicoeur, 1963), and obtained scores from this eigenvector, hereafter called body size, by postmultiplying the $n \times p$ matrix of log-transformed data, where n is the number of observations, by the $p \times 1$ isometric eigenvector. To remove the effects of body size from the log-transformed data, we used Burnaby's method (Burnaby, 1966): we postmultiplied the $n \times p$ matrix of the log-transformed data by a $p \times p$ symmetric matrix, $\mathbf{L}$, defined as:

$$\mathbf{L} = \mathbf{I}_p - \mathbf{V}(\mathbf{V}^T \mathbf{V})^{-1} \mathbf{V}^T$$

where $\mathbf{I}_p$ is a $p \times p$ identity matrix, $\mathbf{V}$ is the isometric size eigenvector defined above, and

$\mathbf{V}^T$ is the transpose of matrix $\mathbf{V}$ (Rohlf and Bookstein, 1987). Hereafter, we refer to the resulting size-adjusted variables as shape variables.

To identify morphometric and meristic variables that best discriminate among species, we used a stepwise discriminant analysis coupled with 100-fold cross-validation to measure classification performance (Quinn and Keough, 2002) using the package *klaR* in R v. 2.12.0 (R Development Core Team, 2009).

RESULTS

Phylogenetic Analyses

We sequenced 1,088 base pairs of the mitochondrial *ND2* gene and adjacent tRNAs, which contained 82 variable sites and 633 parsimony-informative characters. Thirty-nine new mtDNA sequences from 34 localities (Fig. 1) are reported and aligned with 14 previously published sequences.

A comparison of the partitioned Bayesian analyses to the unpartitioned analyses strongly favored the partitioned strategy (Bayes Factors > 860). We observed convergence among multiple Bayesian runs and utilized post-burnin samples (burnin = 1,000) to estimate model parameters and tree topology (Fig. 2). The partitioned ML analysis produced a single tree (Fig. 3, $\ln L = -16,649.1489$) that had a similar topology to the partitioned Bayesian consensus tree at well-supported nodes. The Parsimony analysis produced 54 equally most parsimonious trees (TL = 3,832, CI = 0.337683, RI = 0.682552, RC = 0.230486, HI = 0.662317; Fig. 4). Subspecies formed strongly supported monophyletic groups in all analyses, with the exception of specimens of *A. c. tandai* from Acre. All analyses also recovered a paraphyletic *A. chrysolepis* with regard to *A. bombiceps* and *A. meridionalis* (Figs. 2–4). Sampled individuals of *A. chrysolepis*, *A. bombiceps*, and *A. meridionalis* were members of one of two clades; one (Clade A) composed of *A. c. chrysolepis*, *A. c. tandai*,

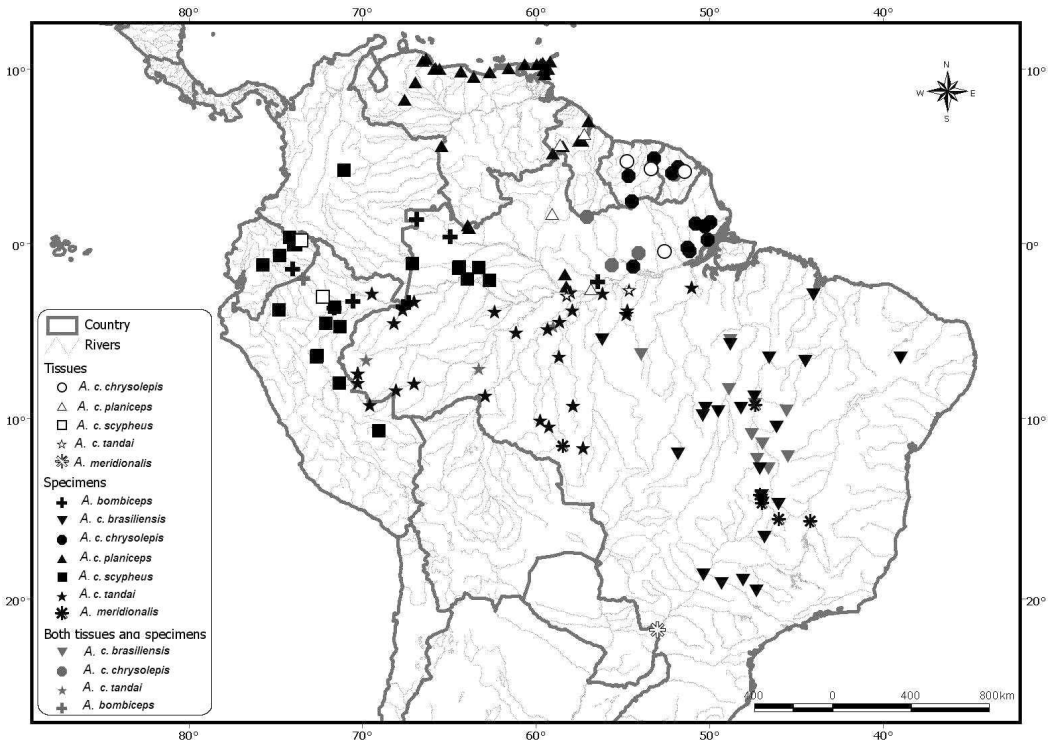


Figure 1. Distribution of material examined of *A. c. chrysolepis*, *A. c. scypheus*, *A. c. tandai*, *A. c. brasiliensis*, *A. c. planiceps*, *A. bombiceps* and *A. meridionalis*. Symbols may represent more than one locality.

and *Anolis meridionalis* and another (Clade B) composed of *A. c. brasiliensis*, *A. c. planiceps*, *A. c. scypheus*, and *A. bombiceps*. Relationships among taxa in clade B were similar across all trees, with an *A. bombiceps* + *A. c. scypheus* clade and an *A. c. planiceps* + *A. c. brasiliensis* clade that are sister taxa. Relationships within Clade A varied depending on the analysis. Parsimony analysis recovered *A. c. tandai* from Acre (LSUMZ H13599) as the sister taxon of the *A. c. tandai* + *A. c. chrysolepis* clade. The ML and Bayesian trees, on the other hand, recovered the Acre *A. c. tandai* as the sister taxon of *A. c. chrysolepis*, but with low bootstrap support. The *A. c. chrysolepis* + *A. c. tandai* clade was well supported in all analyses, whereas the *A. c. chrysolepis* + *A. c. tandai* + *A. meridionalis* clade received poor nodal supported.

Uncorrected pairwise distances among lineages in the *A. chrysolepis* species group

ranged from 5.0% between *A. c. tandai* and *A. c. chrysolepis* to 22.1% between *A. c. scypheus* and *A. meridionalis* (Table 2).

Both the SH and AU tests (Table 3) found that the alternative hypothesis of a monophyletic *A. chrysolepis*, excluding both *A. bombiceps* and *A. meridionalis*, resulted in a significantly worse tree than the ML tree. The ML tree constrained to exclude just *A. meridionalis* was not significantly worse than our best ML tree. Similarly, both tests found no significant difference between a tree constraining a monophyletic *A. c. tandai* and our best ML tree. Bayesian Posterior Probabilities of alternative hypotheses showed little to no support (e.g., Posterior Probabilities < 0.05) for a monophyletic *A. chrysolepis* excluding *A. bombiceps* and *A. meridionalis*, as well as a monophyletic *A. c. tandai*. The Bayesian Posterior Probability of a monophyletic *A. chrysolepis* + *A. bombiceps*, excluding *A. meridionalis*, received moderate support.

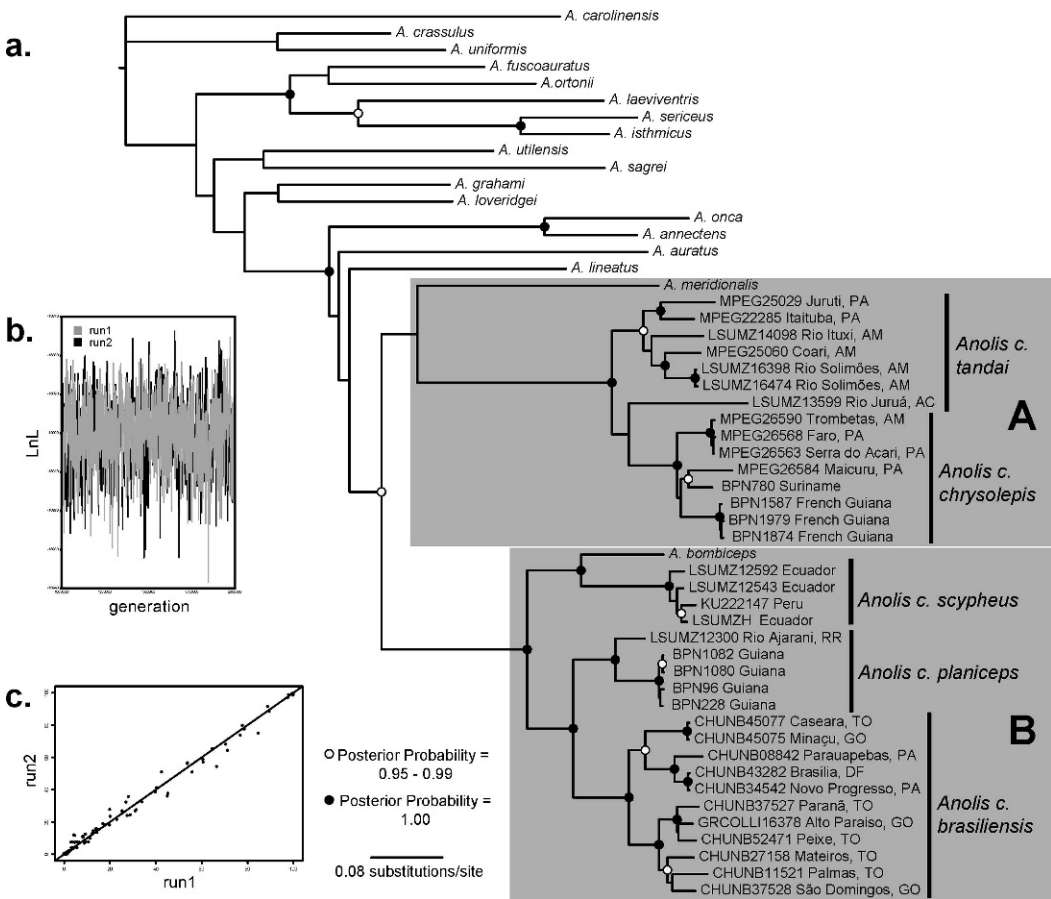


Figure 2. Results of the partitioned Bayesian analysis. a. Phylogeny of the *Anolis chrysolepis* species group and outgroups. Bayesian posterior probabilities >0.95 are indicated by circles at nodes. b. Trace plot of post-burnin log likelihood values for the two Bayesian runs. c. Bivariate plot of the split frequencies for the two Bayesian runs.

Morphological Analyses

The stepwise discriminant analysis applied on body size and all shape variables selected tibia length, interparietal length, and snout-vent length (all size-adjusted) as the most powerful discriminators of *A. chrysolepis* spp., *A. bombiceps*, and *A. meridionalis*, with a classification accuracy of 0.67 based on cross-validation. The first two linear discriminant functions based on these three variables explained about 99% of the total variation, the first function mainly representing a contrast between relative tibia length (–) versus relative SVL (+), and the second function repre-

senting primarily the variation in interparietal length (Table 4, Fig. 5). Results indicate that *A. meridionalis* and *A. c. brasiliensis* have short tibias and elongate bodies relative to total body size, whereas *A. c. tandai* and *A. c. chrysolepis* have long tibias and short bodies relative to total body size, and *A. bombiceps*, *A. c. planiceps*, and *A. c. scypheus* have intermediate values of these variables. Additionally, *A. c. planiceps* has the longest, and *A. c. chrysolepis* the shortest, interparietal relative to its body size. Morphologically, *A. c. chrysolepis* and *A. c. tandai* are very similar, whereas *A. meridionalis* is the most divergent species,

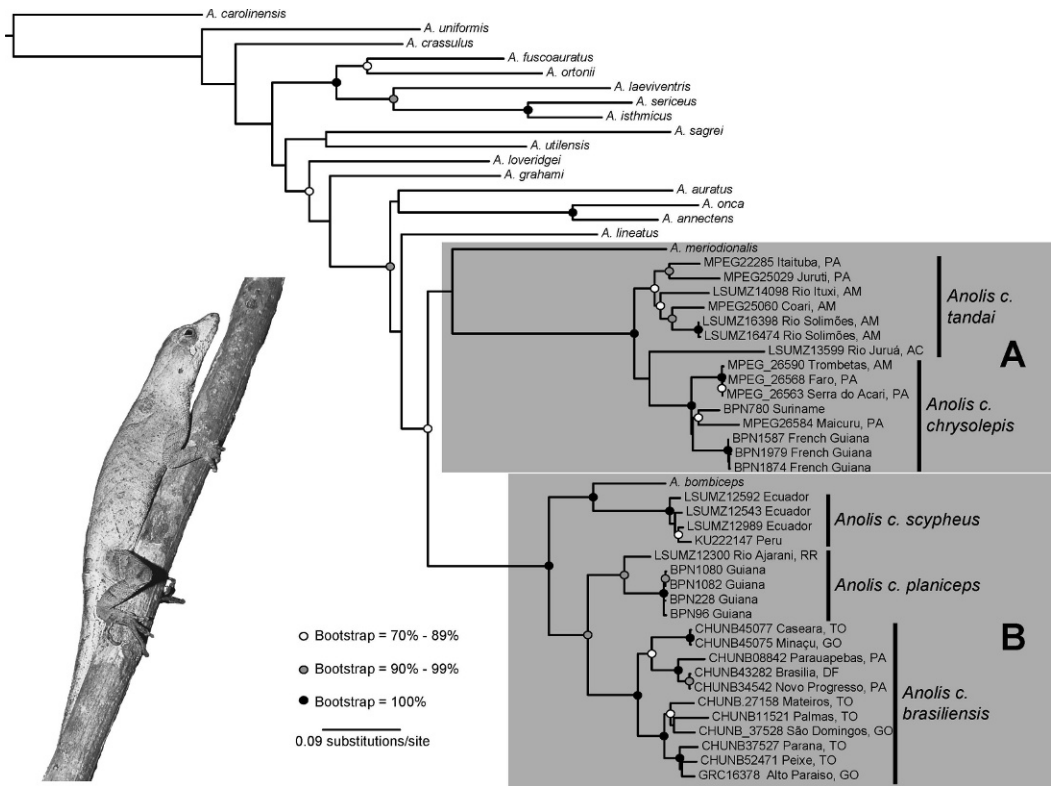


Figure 3. Partitioned Maximum Likelihood phylogeny of the *Anolis chrysolepis* species group and outgroups. Bootstrap values >70% are indicated by circles at nodes. Photo: *Anolis brasiliensis* from São Domingos, Goiás, Brazil. Tony Gamble.

followed by *A. c. brasiliensis*. Nevertheless, classification accuracy based on morphology was moderate.

The stepwise discriminant analysis applied on meristic counts selected canthals, fourth toe lamellae, and scales between second canthals as the most powerful discriminators of the species, with a classification accuracy of 0.83 based on cross-validation (Fig. 6). The first two linear discriminant functions based on these three variables explained about 93% of the total variation. The first function mainly represented a contrast between canthals and scales between second canthals (–) versus fourth toe lamellae (+), whereas the second function primarily represented the variation in fourth toe lamellae and canthals (Table 5, Fig. 6).

Results indicate discrimination 1) in the number of canthals among *A. c. planiceps*

and *A. meridionalis* (small); *A. bombiceps*, *A. c. tandai*, and *A. c. chrysolepis* (large); and *A. c. brasiliensis* and *A. c. scypheus* (intermediate); 2) in the number of fourth toe lamellae among *A. c. chrysolepis* and *A. meridionalis* (small); *A. c. brasiliensis*, *A. c. planiceps*, and *A. c. scypheus* (large); and *A. bombiceps* and *A. c. tandai* (intermediate); and 3) in the number of scales between second canthals among *A. meridionalis* (few), *A. c. scypheus* and *A. tandai* (large), and the remaining species (intermediate). Overall, *A. c. chrysolepis*, *A. c. tandai*, and *A. bombiceps* are more similar, the same happening with *A. c. planiceps*, *A. brasiliensis*, and *A. c. scypheus*. *Anolis meridionalis* is the most divergent species. Classification accuracy based on meristic counts was relatively good.

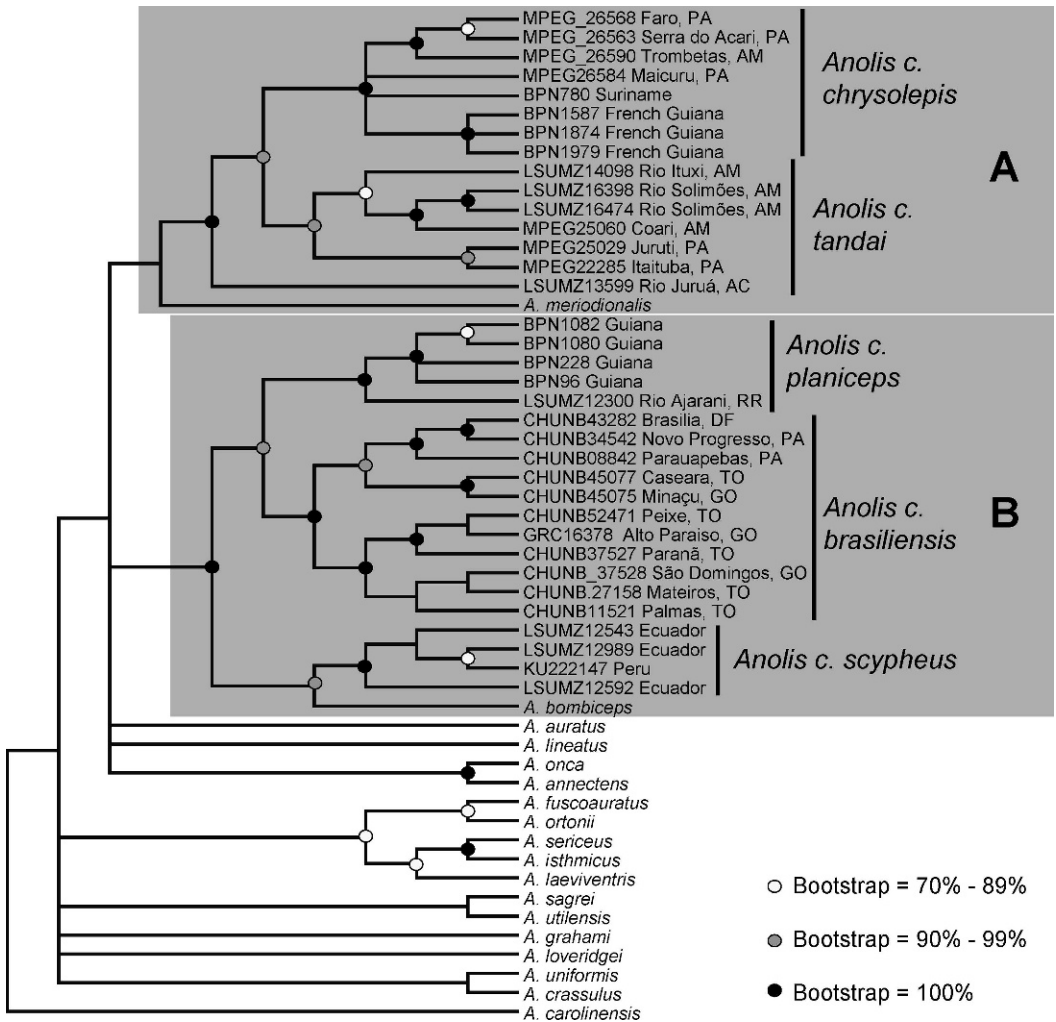


Figure 4. Maximum Parsimony consensus phylogeny of the *Anolis chrysolepis* species group and outgroups.

TABLE 2. NET BETWEEN GROUP DISTANCES FOR *ND2* AMONG THE *ANOLIS CHRYSOLEPIS* GROUP. DISTANCES ABOVE THE DIAGONAL ARE UNCORRECTED *P*-DISTANCES. DISTANCES BELOW THE DIAGONAL WERE MAXIMUM LIKELIHOOD-CORRECTED USING THE GTR MODEL.

	<i>A. c.</i> <i>chrysolepis</i>	<i>A. c.</i> <i>tandai</i>	<i>A. c.</i> <i>planiceps</i>	<i>A. c.</i> <i>brasiliensis</i>	<i>A. c.</i> <i>scyphus</i>	<i>A.</i> <i>bombiceps</i>	<i>A.</i> <i>meridionalis</i>
<i>A. c. chrysolepis</i>	—	0.050	0.195	0.173	0.192	0.189	0.201
<i>A. c. tandai</i>	0.059	—	0.169	0.149	0.167	0.170	0.171
<i>A. c. planiceps</i>	0.274	0.233	—	0.076	0.128	0.136	0.193
<i>A. c. brasiliensis</i>	0.249	0.209	0.093	—	0.114	0.121	0.193
<i>A. c. scyphus</i>	0.270	0.229	0.160	0.155	—	0.104	0.221
<i>A. bombiceps</i>	0.255	0.228	0.156	0.154	0.123	—	0.209
<i>A. meridionalis</i>	0.267	0.232	0.279	0.270	0.298	0.267	—

TABLE 3. COMPARISONS OF MAXIMUM LIKELIHOOD (ML) TREE SCORES ($-\ln L$) AND P VALUES OF THE SH AND AU TESTS BETWEEN OUR BEST ML TREE AND THE CONSTRAINED TREES. BAYESIAN POSTERIOR PROBABILITIES OF ALTERNATIVE HYPOTHESES ARE ALSO SHOWN.

Hypothesis	$-\ln L$	Difference $-\ln L$	SH Test (P)	AU Test (P)	Bayesian Posterior Probability
Optimal tree	-16,900.3198	n/a	n/a	n/a	n/a
Monophyletic <i>A. chrysolepis</i> group	-16,963.9620	-63.6423	<0.001	<0.0001	0.000
Monophyletic <i>A. chrysolepis</i> group + <i>A. bombiceps</i>	-16,900.9392	-0.6194	0.867	0.571	0.219
Monophyletic <i>A. c. tandai</i>	-16,903.4524	-3.1326	0.726	0.420	0.020

DISCUSSION

The molecular phylogenetic analyses recovered six species-level taxa as part of the *A. chrysolepis* species group. These taxa can also be morphologically distinguished on the basis of morphometric and meristic characters. Even though we cannot infer relationships among these taxa on the basis of the meristic discriminant analysis, the results of this analysis are consistent with the existence of two clades: one containing *A. c. tandai*, *A. c. chrysolepis*, and *A. bombiceps* and another clade containing *A. c. brasiliensis* and *A. c. planiceps*. Meristic characters in *A. c. scypheus* appear to be intermediate between these two groups, which is also consistent with it being (together with *A. bombiceps*) the sister clade to *A. c. brasiliensis* + *A. c. planiceps*. *Anolis meridionalis* was quite distinct from other members of the *A. chrysolepis* species group on the basis of meristic characters.

We define the *A. chrysolepis* species group as the clade originating with the most

recent common ancestor of *A. c. chrysolepis* and *A. c. brasiliensis*. *Anolis meridionalis* has not historically been allied with the *A. chrysolepis* species group because of its unique morphology. In particular, *A. meridionalis* differs from other members of the *A. chrysolepis* species group by having digital dilatations on phalanx ii and iii continuous with scales under phalanx i, instead of forming the prominent border observed in the *A. chrysolepis* subspecies and *A. bombiceps*. Although the node leading to the *A. chrysolepis* species group, including *A. meridionalis*, was well supported in the ML and Bayesian analyses, the presence of *A. meridionalis* in clade A received poor support in all phylogenetic analyses. For this reason, we could not reject the alternative hypothesis of a monophyletic *A. chrysolepis* group exclusive of *A. meridionalis*. This means that inclusion of *A. meridionalis* in the *A. chrysolepis* species group is still uncertain. Future phylogenetic analyses that include additional *A. meridio-*

TABLE 4. LINEAR DISCRIMINANT ANALYSIS OF THREE MORPHOMETRIC VARIABLES THAT BEST DISTINGUISH THE SPECIES AND SUBSPECIES OF *ANOLIS* STUDIED. VALUES REPRESENT MEANS OF SCALED, SIZE-ADJUSTED VARIABLES FOR EACH SPECIES AND COEFFICIENTS OF VARIABLES ON FIRST AND SECOND LINEAR DISCRIMINANT FUNCTIONS (LDF 1, LDF 2). PROPORTION OF TOTAL VARIATION EXPLAINED BY EACH LDF IN PARENTHESES.

<i>Anolis</i> Species	Tibia Length	Interparietal Length	Snout-Vent Length
<i>A. bombiceps</i>	0.57	-0.35	0.78
<i>A. c. brasiliensis</i>	-0.91	0.46	0.15
<i>A. c. chrysolepis</i>	0.67	-1.02	0.14
<i>A. meridionalis</i>	-1.67	0.58	1.46
<i>A. c. planiceps</i>	-0.02	0.79	-0.55
<i>A. c. scypheus</i>	-0.05	-0.32	-0.19
<i>A. c. tandai</i>	0.87	-0.60	0.11
LDF 1 (0.86)	-1.72	0.53	1.05
LDF 2 (0.13)	-0.60	-1.11	0.49

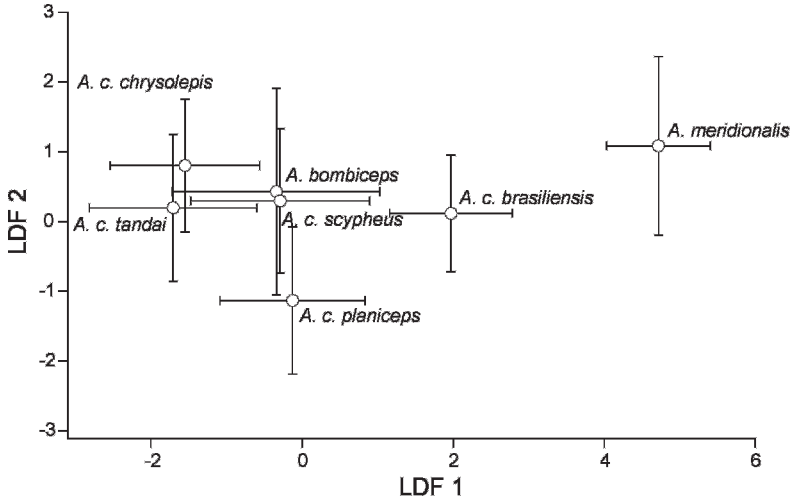


Figure 5. Means (open circles) and standard deviations (error bars) of scores on first (LDF 1) and second (LDF 2) linear discriminant functions of tibia length, interparietal length, and snout-vent length (all size-adjusted; see text for details) for seven subspecies and two species of *Anolis*.

nalis samples and data from nuclear loci may help resolve this issue.

All described taxa in the molecular analyses formed well-supported, monophyletic groups, with the exception of *A. c. tandai*. The *A. c. tandai* individual from Acre fit the morphological diagnosis we present in this study but was either the sister taxon to *A. c. chrysolepis* (ML and Bayesian analyses) or the sister taxon to the

A. c. chrysolepis + *A. c. tandai* clade (parsimony analysis). The apparent paraphyly of *A. c. tandai* may be due to several phenomena, none of which are mutually exclusive. One possibility is phylogenetic error due to incomplete taxonomic sampling or lack of data (Graybeal, 1998; Mitchell et al., 2000). It is also possible that individuals from the Acre population represent an as yet undescribed, morphologically cryptic

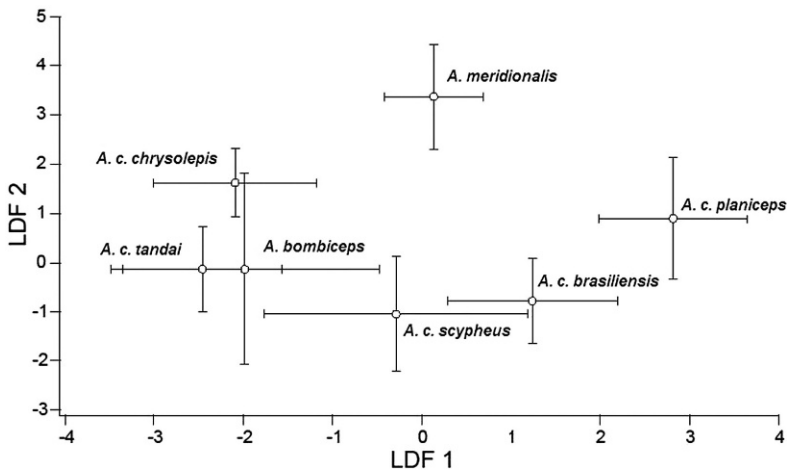


Figure 6. Means (open circles) and standard deviations (error bars) of scores on first (LDF 1) and second (LDF 2) linear discriminant functions of canthals, fourth toe lamellae, and scales between second canthals for seven species of *Anolis*.

TABLE 5. LINEAR DISCRIMINANT ANALYSIS OF THREE MERISTIC COUNTS THAT BEST DISTINGUISH THE SPECIES AND SUBSPECIES OF *ANOLIS* STUDIED. VALUES REPRESENT MEANS OF SCALED VARIABLES FOR EACH SPECIES AND COEFFICIENTS OF VARIABLES ON FIRST AND SECOND LINEAR DISCRIMINANT FUNCTIONS (LDF 1, LDF 2). PROPORTION OF TOTAL VARIATION EXPLAINED BY EACH LDF IN PARENTHESES.

<i>Anolis</i> Species	Canthals	Fourth Toe Lamellae	Scales Between Second Canthals
<i>A. bombiceps</i>	1.13	-0.63	-.022
<i>A. c. brasiliensis</i>	-0.09	0.72	-0.67
<i>A. c. chrysolepis</i>	0.41	-1.50	0.21
<i>A. meridionalis</i>	-0.77	-1.80	-1.05
<i>A. c. planiceps</i>	-1.48	0.42	-0.56
<i>A. c. scyphus</i>	0.09	0.56	0.99
<i>A. c. tandai</i>	0.96	-0.66	0.75
LDF 1 (0.74)	-1.38	0.92	-0.69
LDF 2 (0.18)	-0.92	-1.36	-0.20

species. Incomplete lineage sorting can also result in discordance between individual gene trees and the species tree because of the retention and/or sorting of ancestral polymorphisms, particularly when populations have diverged recently, have a large effective population size, or both (Maddison, 1997; Ballard and Whitlock, 2004; Maddison and Knowles, 2006). Additional phylogenetic analyses incorporating nuclear genes and additional taxa, as well as using methods that incorporate coalescent processes and incomplete lineage sorting, would be useful in clarifying relationships among *A. c. tandai* populations.

Our results show broad congruence among molecular and morphological data sets that are consistent with independent evolutionary lineages. Most importantly, each of these lineages is morphologically diagnosable. Genetic distances among sister taxa in the *A. chrysolepis* group were also comparable to ND2 distances among sister species in other squamate taxa (Macey et al., 1998, 1999; Glor et al., 2001; Oliver et al., 2009). Therefore, we elevate each subspecies to species status under the general lineage species concept (de Queiroz, 1998, 1999, 2005, 2005a, 2005b, 2007). To facilitate future studies, each species, including *A. bombiceps* and *A. meridionalis*, is diagnosed below and an identification key is provided, considering morphological data collected for this study as well as data from the literature. Table 6

compares the main meristic and morphometric characters.

Taxonomy/Species Accounts

All descriptions of color pattern are based on literature, photographs of live animals, and preserved specimens.

Anolis chrysolepis Duméril and Bibron, 1837.

Anolis chrysolepis Duméril and Bibron, 1837:94 (lectotype MHNP 2456, type locality: La Mana, French Guiana); Cunha, 1961:60; Peters and Donoso-Barros, 1970:61; Avila Pires et al., 2010:94.

Anolis chrysolepis chrysolepis; Vanzolini and Williams, 1970:85; Hoogmoed, 1973:112; Hoogmoed and Avila-Pires, 1989:168.

Norops nitens chrysolepis; Savage and Guyer, 1991:366.

Anolis nitens chrysolepis; Avila-Pires, 1995:75.

Abbreviated Description. Maximum SVL 74 mm. Vertebral region with distinctly enlarged scales, middorsal row largest; number of rows of enlarged scales increases posteriorly. Scales on upper arms smaller than, to subequal to, vertebral scales. Supraorbital semicircles with scarcely enlarged scales. Supraocular scales keeled, slightly larger than or subequal to scales

TABLE 6. COMPARISONS OF MERISTIC CHARACTERS, BODY PROPORTIONS AND MEASUREMENTS (IN MILLIMETERS) AMONG THE SPECIES OF MEMBERS OF THE *ANOLIS CHRYSOLEPIS* GROUP.

Character*	A. <i>chrysolepis</i>	A. <i>tandai</i>	A. <i>scypheus</i>	A. <i>planiceps</i>	A. <i>brasiliensis</i>	A. <i>bombiceps</i>	A. <i>meridionalis</i>
No. of specimens	50	96	49	72	116	11	9
max. svl (mm)	74	70	80	76	69	74	56
midbody	101–156	91–150	110–174	110–149	112–155	125–137	79–106
midbody median and standard deviation values	123.18 ± 12.8	134.06 ± 7.43	145.94 ± 14.59	125.84 ± 9.46	129.12 ± 10.78	131.37 ± 3.08	92.22 ± 9.39
slabials	11–15	10–14	10–14	10–14	10–13	10–12	9–11
ilabials	11–15	10–14	10–13	9–11	8–12	10–13	8–11
prostrals	4–7	5–9	5–7	4–7	5–7	5–6	5
loreal	5–10	5–9	7–10	6–9	6–10	6–7	6–7
canthals	6–10	6–11	6–11	6–8	5–9	9–11	5–6
scales bet 2 ^a canthals	9–14	10–15	9–15	8–14	8–14	9–12	8–10
scales bet semicircsorbit	1–4	1–4	1–5	1–3	0–3	1–3	0–1
interp-semicircsorbit	1–5	2–4	2–5	1–3	1–4	2–4	0–2
pmentals	4–8	4–8	4–8	4–8	4–8	6–8	5–8
lam-4fg	12–17	14–19	15–21	14–20	15–20	15–19	10–12
lam-4toe	18–26	20–28	26–35	19–34	25–32	21–29	18–24
tail/svl	1.15–2.17	1.48–2.42	1.66–2.53	1.33–3.27	1.22–2.54	1.69–2.18	1.97–2.38
mouth/svl	0.13–0.25	0.19–0.24	0.18–0.24	0.19–0.26	0.18–0.26	0.19–0.21	0.19–0.21
interp/head-w	0.07–0.19	0.08–0.24	0.11–0.25	0.16–0.32	0.11–0.30	0.10–0.28	0.16–0.30
tibia/svl	0.20–0.38	0.30–0.43	0.26–0.38	0.27–0.36	0.26–0.34	0.29–0.36	0.22–0.27
Interparietal	0.64–1.65	0.84–2.22	0.81–2.37	0.98–2.79	0.92–2.65	1.1–2.9	1.28–2.07
head-w	5.93–10.5	5.31–11.41	5.67–13.08	4.95–12.14	7.25–11.75	8.96–10.73	6.95–8.07
head-alt	5.00–8.39	4.33–9.65	4.23–11.02	3.28–9.39	5.56–9.2	7.62–8.33	4.88–6.36
orbdist	4.01–6.39	3.4–7.19	3.04–8.31	3.31–8.41	4.39–7.26	6.09–6.65	4.02–4.86
cardiam	0.40–1.41	0.75–1.89	0.40–2.23	0.47–1.71	0.87–2.00	0.88–1.68	0.84–1.09
nostrilsdis	1.25–2.37	1.14–2.7	1.10–2.95	1.07–2.76	1.40–2.82	2.0–2.36	1.42–1.73
mouth to ear	1.13–3.03	1.6–2.92	1.09–3.09	1.06–3.52	0.96–2.72	1.47–2.18	1.28–1.53
snout	2.97–6.69	3.35–7.00	2.78–7.72	2.69–7.17	3.92–8.14	4.67–6.46	4.8–5.82
max. toe IV	20.94	24.42	26.91	28.31	21.63	25.51	14.12
max. foot	25.63	30.24	31.46	31.89	26.48	22.66	17.39
toe IV width	0.55–1.30	0.47–1.29	0.61–1.48	0.51–1.46	0.92–1.65	0.96–1.26	0.61–0.91

* Abbreviations: max. svl = maximum snout-vent length; midbody = number of scales around midbody; slabials = total number of supralabials; ilabials = total number of infralabials; prostrals = total number of postrostrals; scales bet 2^a canthals = number of scales on the snout between the second canthals; scales bet semicircsorbit = minimum number of scales between supraorbital semicircles; interp-semicircsorbit = minimum number of scales between any of the supraorbital semicircles and interparietal; pmentals = number of postmentals; lam-4fg = number of expanded lamellae under the fourth finger; lam-4toe = number of expanded lamellae under fourth toe; tail/svl, mouth/svl, tibia/svl = respectively, the rates of the tail, mouth, and tibia length with the snout-vent length; interp/head-w = the rate of interparietal width with head width; interparietal = interparietal width; head-w = head width; head-alt = head height; orbdist = minimum distance between orbits; cardiam = ear diameter; nostrilsdis = minimum distance between nostrils; mouth to ear = minimum distance between mouth and ear; snout = from the tip of snout to anterior margin of orbit; max. toe IV = from toe IV base to the heel; max. foot = fourth toe length from toe nail to toe base; toe IV width = fourth toe width.

on snout, grading into granules laterally and posteriorly. Interparietal subequal to or slightly larger than adjacent scales (Fig. 7A, B).

Color in Preservative. Color pattern sexually dimorphic. Male dorsal color pale or grayish-brown with or without a wide light vertebral band bordered by a grayish-brown irregular band laterally. Paired triangular spots may be present along back; most specimens with paired triangular spots on sacral region. Female dorsal pattern less variable. A thin dark brown line begins at

posterior corner of each eye at each side, converging toward neck and continuing along the body, where they delimit a lighter or plumbeous vertebral band that darkens and expands laterally on tail.

Dewlap Color in Life and Preservative. In preservative, male dewlap skin royal blue, dark blue, or blackish, with light scales or blue scales toward rim. Female dewlap cream, similar to surrounding area; scales may be darker at edge. In life, male dewlap skin usually royal blue or blackish blue with



Figure 7. *Anolis chrysolepis* species group. A) *Anolis chrysolepis* female from Nassau Plateau, Suriname (Photo: Robert Langstroth), B) *Anolis chrysolepis* male from Faro, Pará, Brazil (Photo: Waldima Rocha), C) *Anolis tandai* female from Rio Juruá, Acre (Photo: Laurie J. Vitt), D) *Anolis tandai* male from Rio Juruá, Acre, Brazil (Photo: Laurie J. Vitt), E) *Anolis tandai* female from Rio Ituxi, Amazonas, Brazil (Photo: Laurie J. Vitt), F) *Anolis tandai* male from Amazonas, Brazil (Photo: Laurie J. Vitt).

light scales or blue scales along rim. Avila-Pires (1995) mentioned a cobalt-blue juvenile male dewlap (RMNH 24673) with white to orange scales, surrounded by a spectrum-orange area that extended through most of ventral surface of head. Female dewlap skin usually yellowish to orange with gray or cream scales; an orange

lateral area extended through most of ventral surface of head may be present. Hoogmoed and Avila-Pires (1991) mentioned a female from French Guiana with yellow dewlap with orange scales, presenting a bluish area toward the rim.

Comparison with Other Species from the A. chrysolepis Species Group. This species

has proportionally the smallest interparietal length among the species of the group (Table 6). It differs from its sister taxon *A. tandai* mainly by a lower number of postrostral scales (4–7 in *A. chrysolepis* and 5–9 in *A. tandai*) and by female dewlap color, generally cream in *A. chrysolepis*; cream with a large central blue spot in *A. tandai*.

Distribution. Southern Guyana, Suriname, French Guiana and northern Brazil, in the states of Amapá and Pará.

Anolis tandai Avila-Pires, 1995.

Anolis chrysolepis; Vanzolini, 1986:18; Gascon and Pereira, 1993:181.

Anolis nitens tandai Avila-Pires, 1995:80 (holotype MPEG 15850, type locality: Rio Urucu, Amazonas state, Brazil); Icochea et al., 2001:140; Vitt et al., 2001:401; Santos-Jr et al., 2007:9; Avila-Pires et al., 2009:116.

Abbreviated Description. Maximum SVL 70 mm. Vertebral region with slightly enlarged scales; number of rows of enlarged scales increases posteriad. Scales on upper arms smaller than, or subequal to, vertebral scales. Supraorbital semicircles with scarcely enlarged scales. Supraocular scales weakly to distinctly keeled, approximately same size as scales on snout, laterally and posteriorly grading into granules, anteriorly surrounded by smaller scales. Interparietal moderately small, larger than adjacent scales (Fig. 7C–F).

Color in Preservative. Color pattern sexually dimorphic. In males, vertebral region usually distinct from flanks, with unclear limits between these areas. A pair of subtriangular dark spots present on sacral region. Some specimens may present sinuous lines, assuming subtriangular shapes along dorsum. Females usually with a well-delimited vertebral band, similar to *Anolis chrysolepis* females; occasionally dorsal pattern similar to males.

Dewlap Color in Life and Preservative. In preservative, male dewlap royal blue or blackish-blue with light scales. Female dewlap with central blue spot surrounded

by a cream area; scales usually light colored. In life, male dewlap skin frequently blue or blackish, with light scales. Avila-Pires (1995) mentioned the dewlap in MPEG 15986 as “ultramarine with cream-color scales on rim.” Dewlap in females, when extended, presents a large and central blue spot, surrounded by a cream area. Scales are frequently cream to orange. When not extended, dewlap presents a light rim and is blue laterally. Avila-Pires (1995) described the holotype MPEG 15850 female dewlap color as “sulphur-yellow with a large indigo-blue spot.”

Comparison with Other Species from the A. chrysolepis Species Group. As already mentioned by Avila-Pires (1995), this species has the longest tibia in relation to SVL (0.30–0.43). For differences with *A. chrysolepis*, see above. Avila-Pires (1995) also mentioned the possible sympatry with *A. bombiceps*, which also has a blue or blackish blue dewlap (with no sexual dimorphism), but they can be distinguished by female dewlap color (a central blue spot, surrounded by a pale area in *A. tandai*), by the minimum number of scales between supra-orbital semicircles (1–4 in *A. tandai* and 1–2 in *A. bombiceps*) and by the number of postmentals (4–8 in *A. tandai* and 6–8 in *A. bombiceps*).

Distribution. South of the Amazon River and west of the Tapajós River, in Brazil (states of Pará, Amazonas, Rondônia, Acre, and north of Mato Grosso), and in Peru.

Anolis planiceps Troschel, 1848.

Anolis planiceps Troschel, 1848:649 (holotype ZMB 529, type locality: Caracas, Venezuela).

Anolis chrysolepis planiceps; Vanzolini and Williams, 1970:85; Hoogmoed, 1973:125; Myers and Donnelly, 2008:100.

Anolis chrysolepis; Beebe, 1944:97; O’Shea, 1989:69; Zimmerman and Rodrigues, 1990:449; Martins, 1991:182.

Anolis eewi Roze, 1958:311 (holotype FMNH 74040, type locality: Chimantapui, Bolívar, Venezuela).

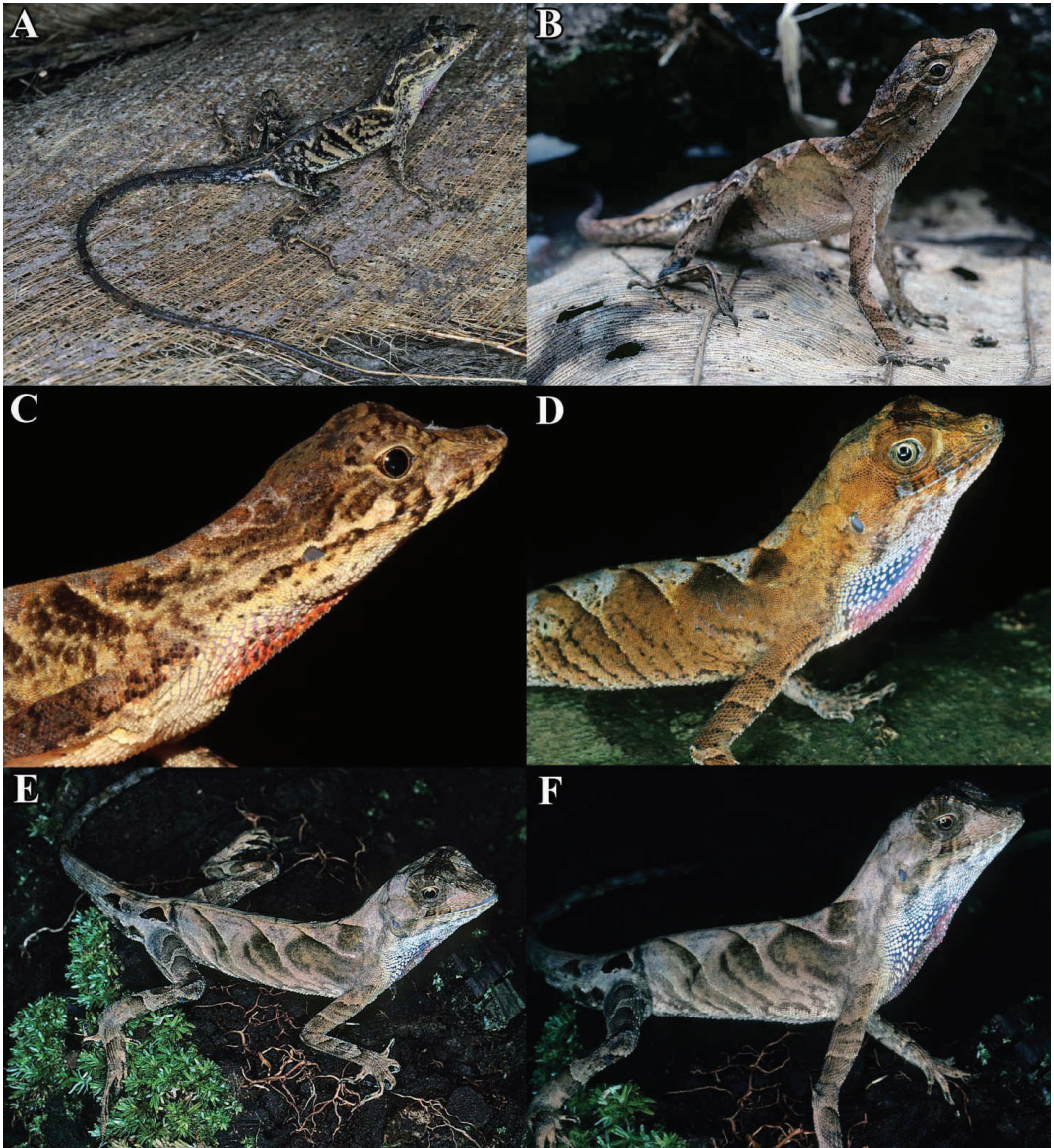


Figure 8. *Anolis chrysolepis* species group. A) *Anolis planiceps* from Roraima, Brazil (Photo: Laurie J. Vitt), B) *Anolis planiceps* from Guatopo, Venezuela (Photo: Laurie J. Vitt), C) *Anolis planiceps* from Cuyuni-Mazaruni, Guyana (Photo: Robert Langstroth), D) *Anolis scypheus* from Ecuador (Photo: Laurie J. Vitt), E) *Anolis scypheus* from Ecuador (Photo: Laurie J. Vitt), F) *Anolis scypheus* from Ecuador (Photo: Laurie J. Vitt).

Anolis nitens: Boulenger, 1885:91; Beebe, 1944:200.

Norops nitens nitens; Savage and Guyer, 1991:366.

Anolis nitens nitens; Avila-Pires, 1995:70; Vitt et al., 2008:84.

Abbreviated Description. Maximum SVL 76 mm. Double row of enlarged vertebral scales extending from nape to base of tail; few to several rows of weakly keeled scales, increasing in number caudally, forming a gradual transition between double row of enlarged scales and

granules on flanks. Scales of upper arms markedly larger than vertebral scales. Supraorbital semicircles with enlarged scales, forming pronounced ridge in some specimens. Supraocular region with distinct group of enlarged, weakly keeled, scales surrounded by smaller scales. Interparietal distinctly larger than adjacent scales (Fig. 8A–C).

Color in Preservative. No sexual dimorphism in color pattern. Specimens usually have many chevrons along back, with tips directed posteriorly, sometimes forming the posterior border of rhomboid figures. A pair of triangular spots commonly present on sacral region. Myers and Donnelly (2008) described the color pattern of two adult males and one adult female as “orange with white or grayish white scales in basal rows, scales darker gray or blackish gray in distal rows.”

Dewlap Color in Life and Preservative. Dewlap red, fading rapidly in preserved specimens, appearing cream-white, with light scales. A lateral lavender area may be present as mentioned by Avila-Pires (1995). In life, dewlap skin orange to reddish with grayish to cream scales. Myers and Donnelly (2008) found variation in the dewlap of four juveniles, including a female that had “a large bluish black basal spot on the dewlap, which had a bright orange periphery and mostly white scales (only a few dark scales).”

Comparison with the Other Species from the A. chrysolepis Species Group. This species has the proportionately largest interparietal scale. It differs from its sister taxon *A. brasiliensis* mainly by dewlap color (red in *A. planiceps* and blue or grayish/blackish blue in *A. brasiliensis*) and body size (*A. planiceps* reaches 76 mm, whereas *A. brasiliensis* reaches 69 mm).

Distribution. Venezuela, Trinidad, Guyana, and the states of Roraima and Amazonas on the northern part of Brazil.

Anolis brasiliensis Vanzolini and Williams, 1970.

Anolis chrysolepis; Amaral, 1937:1722.

Anolis chrysolepis brasiliensis; Vanzolini and Williams, 1970:85 (holotype MZUSP

10319, type locality Barra do Tapirapés, Mato Grosso, Brazil); Williams and Vanzolini, 1980:99; Vanzolini, 1981:253, 1986:3; Cunha et al., 1985:23.

Norops nitens brasiliensis; Savage and Guyer, 1991:366.

Anolis nitens brasiliensis; Avila-Pires, 1995:70; Werneck and Colli, 2006:1987.

Abbreviated Description. Maximum SVL 69 mm. Double row of enlarged vertebral scales from nape to base of tail; few to several rows of dorsal scales with weak keels, increasing in number caudally, gradually transitioning between double row of enlarged scales and granules on flanks. Scales of upper arms markedly larger than vertebral scales. Scales on snout from moderately keeled to smooth, heterogeneous in size, with no distinction between anterior and posterior scales. Supraorbital semicircles with enlarged, generally smooth scales. Supraocular region with most scales large and weakly keeled, surrounded by small scales. Interparietal distinctly larger than adjacent scales (Fig. 9A–D).

Color in Preservative. No sexual dimorphism in color pattern. Dorsal color grayish-brown or pale white, either uniform or not. A light vertebral band may be present, either narrow with undefined margins or wide; in both cases surrounded by darker area. A pair of triangular spots on sacral region commonly present, may be accompanied by second pair at the base of tail. Ventral region usually pale-white, may be marbled with brown spots.

Dewlap Color in Life and Preservative. Dewlap blue or grayish-blue, with light or grayish scales. In life, dewlap usually grayish blue or blackish blue, with dark scales varying from light-cream to dark gray. Some specimens from Tocantins state show the dewlap skin grayish-green tending to yellowish-beige along rim, with scales grayish-brown or pale-cream tending to brownish along rim. Some irregular light-blue lines may be present (Fig. 8D). Vanzolini and Williams (1970) do not describe the dewlap color but refer to the frontispiece plate

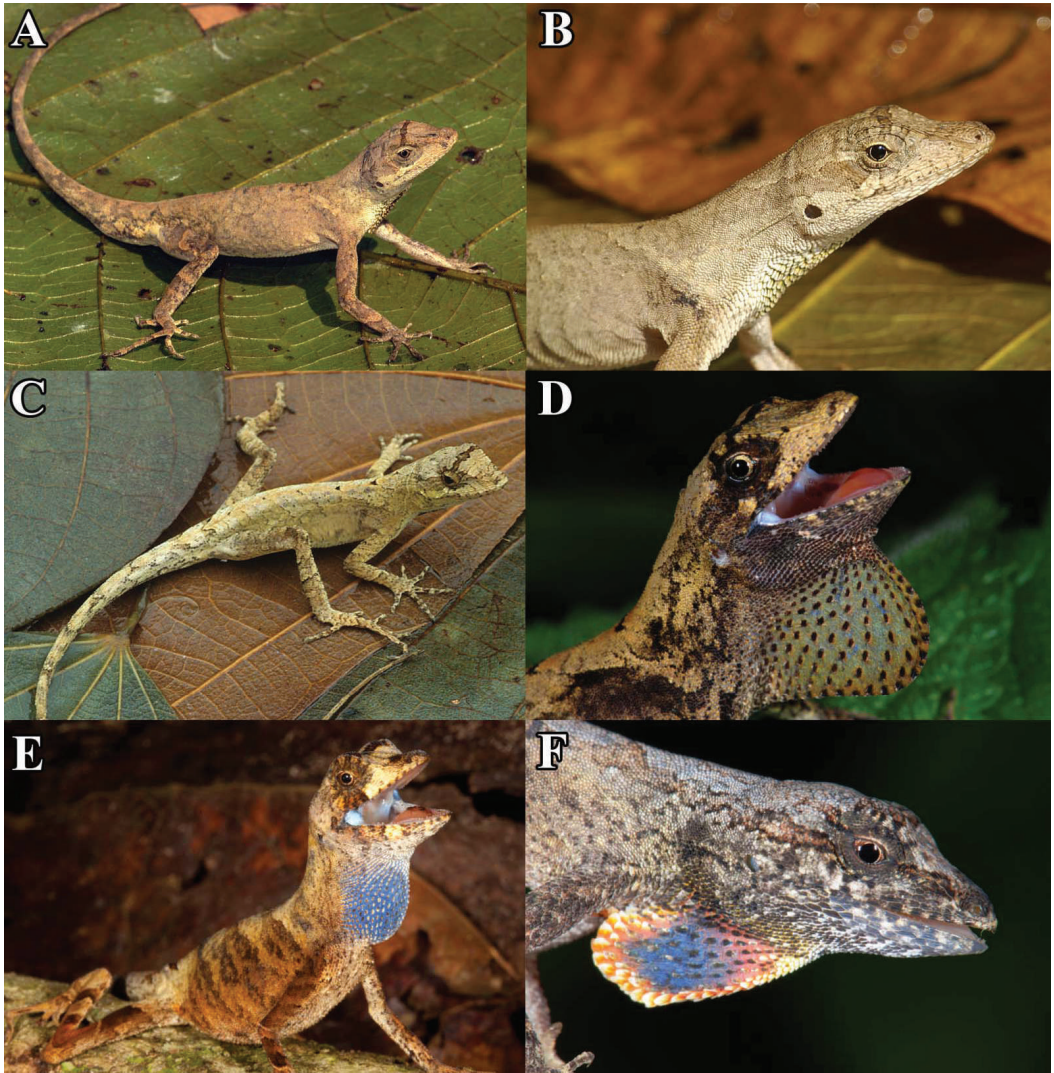


Figure 9. *Anolis chrysolepis* species group. A) *Anolis brasiliensis* from Cantão, Tocantins, Brazil (Photo: Laurie J. Vitt), B) *Anolis brasiliensis* from Cantão, Tocantins, Brazil (Photo: Laurie J. Vitt), C) *Anolis brasiliensis* from Barra do Ouro, Tocantins, Brazil (Photo: Itamar Toniai), D) *Anolis brasiliensis* from Jalapão, Tocantins, Brazil (Photo: Laurie J. Vitt), E) *Anolis bombiceps* from Peru (Photo: Young Cage), F) *Anolis meridionalis* from Tocantins, Brazil (Photo: Itamar Toniai).

representing the dewlap color in life of a male as green with a brown edge along rim. Avila-Pires (1995) observed in specimens from Carajás, Southern Pará, “a blue dewlap, lighter in females, with scales varying from light to dark gray or cream and the surrounding area may be chrome-orange.”

Comparison with the Other Species from the A. chrysolepis Species Group. *Anolis*

brasiliensis, along with *A. bombiceps*, has the largest toe IV among the other species of the *A. chrysolepis* group. *Anolis brasiliensis* differs from *A. planiceps*, mainly by dewlap color (red in *A. planiceps* and blue or grayish/blackish blue in *A. brasiliensis*) and body size (*A. planiceps* reaches 76 mm, whereas *A. brasiliensis* reaches 69 mm). *Anolis brasiliensis* is broadly sympatric with

A. meridionalis, although they occur in different habitats (see *A. meridionalis* description below) and they differ mainly by digital dilatations under phalanx ii and iii that form a prominent border in *A. brasiliensis*.

Distribution. Brazil, in southern Pará, Tocantins, Piauí, Maranhão, Ceará, Goiás, Mato Grosso, Minas Gerais, São Paulo, and Distrito Federal.

Two individuals housed in MCZ under the numbers R-60580 and R-60581 are labeled as paratypes of *A. c. brasiliensis* but have as localities Rio Juruá, Brazil, and Loreto, Peru, respectively. Vanzolini and Williams (1970) did not mention those individuals, which mean that they are not paratypes. Moreover, even though they have typical *A. brasiliensis* characteristics, it is extremely unlikely that the species occurs in these localities. Given their questionable data, we did not consider these individuals in the morphological analyses or elsewhere in this study.

Anolis scypheus Cope, 1864.

Anolis chrysolepis; Guichenot, 1855:15.

Anolis scypheus Cope, 1864:172 (holotype BM 1946.8.855, type locality: "Caracas" according to Boulenger, 1885, but considered an error by Vanzolini and Williams, 1970:85); Boulenger, 1885:90; Goeldi, 1902:16, 32; Cunha, 1961:67; Peters and Donoso-Barros, 1970:66.

Anolis incompertus incompertus Barbour, 1932:99 (holotype MCZ 32309, type locality: Villavicencio, Meta, Colombia).

Anolis chrysolepis scypheus: Vanzolini and Williams, 1970:85; Vanzolini, 1986:3.

Norops nitens scypheus; Savage and Guyer, 1991:366.

Anolis nitens scypheus; Avila-Pires, 1995:78.

Abbreviated Description. Maximum SVL 80 mm. Vertebral scales forming double row of enlarged dorsals along back. Scales of upper arms small relative to other species, larger than vertebral scales. Scales on snout

relatively small, with raised surface. Supra-orbital semicircles with enlarged scales, generally forming pronounced ridge. A small group of enlarged supraocular scales, grading into granules posteriorly and laterally, anteriorly surrounded by smaller scales. Interparietal distinctly larger than surrounded scales (Fig. 8D–F).

Color in Preservative. No sexual dimorphism in color pattern. Dorsum usually with caudally directed chevrons that may form the posterior border of rhomboid figures, similar to the pattern described for *A. planiceps*; or, it may show a broad band with lateral expansions (narrower at nape, extending caudally). A pair of subtriangular dark spots may be present on sacral region.

Dewlap Color in Life and Preservative. Dewlap color in preservative usually pale along rim with pale-cream or blackish scales, and blue with light scales in the center (Fig. 7). In life, dewlap skin red along rim (the red color vanishes very easily in preserved specimens) with red or blackish scales, and blue with pale-cream or light-brown scales laterally. Avila-Pires (1995) described the dewlap color of RMNH 24653 as "cobalt-blue with red rim, scales white with orange center."

Comparison with the Other Species from the A. chrysolepis Species Group. *Anolis c. scypheus* presents the proportionally maximum values of head width, head height, ear diameter, minimum distance between nostrils and SVL among all species of the *A. chrysolepis* species group. For differences with *A. bombiceps*, see *A. bombiceps* diagnosis below.

Distribution. Amazonian Colombia, Ecuador, Peru, and the northwestern part of Amazonas state in Brazil.

Anolis bombiceps Cope, 1876.

Anolis bombiceps Cope, 1876:168 (type apparently lost, type locality: Nauta, Peru); Goeldi, 1902:16, 32; Peters and Donoso-Barros, 1970:49; Vanzolini and Williams, 1970:86; 1986:28; Avila-Pires, 1995:54.

Norops bombiceps; Savage and Guyer, 1989:110.

Abbreviated Description. Maximum SVL 74 mm. Vertebral scales not or only slightly enlarged. Scales on upper arm subequal to, or slightly larger than, vertebral scales. Scales on snout anteriorly small, weakly to distinctly keeled, posteriorly larger, flat, usually uni- or multicarinated. Supraorbital semicircles with enlarged, keeled scales, forming pronounced ridge. Supraocular region with a group of distinctly enlarged scales surrounded posteriorly and laterally by granules. Interparietal scale distinctly larger than adjacent scales (Fig. 9E).

Color in Preservative. No sexual dimorphism in color pattern. Dorsal color usually brown, with irregular dark spots between hind limbs and irregular figures across limbs. V-shaped lines along back, with apex directed posteriorly, may be present.

Dewlap Color in Life and Preservative. Dewlap deep blue or blackish in preservative, with light or dark scales. In life, dewlap skin deep blue with light or dark scales.

Comparisons with the Other Species from the *A. chrysolepis* Species Group. *Anolis bombiceps* is sympatric with *A. scypheus* and may be sympatric with *A. tandai*, of which it can be distinguished by dewlap color (deep blue or blackish in *A. bombiceps*, blue on central region and red along rim in *A. scypheus*, and blue on central region and cream along rim on females of *A. tandai*; males of *A. tandai* and *A. bombiceps* have similar dewlap colors), number of loreals (6–7 in *A. bombiceps*, 5–9 in *A. tandai*, and 7–10 in *A. scypheus*), number of scales between the second canthals (9–12 in *A. bombiceps*, 10–15 in *A. tandai*, and 9–15 in *A. scypheus*), number of scales between supraorbital semicircles (1–2 in *A. bombiceps*, 1–4 in *A. tandai*, and 1–5 in *A. scypheus*) and postmentals (6–8 in *A. bombiceps* and 4–8 in *A. tandai* and *A. scypheus*).

Distribution. Amazonian Colombia, Ecuador, and Peru and in the State of

Amazonas, Brazil. In this study, we reported *A. bombiceps* from two Brazilian localities, both in Amazonas state: Apuí, very close to the Colombian border and São Gabriel da Cachoeira.

Anolis meridionalis Boettger, 1885.

Anolis meridionalis Boettger, 1885a:437 (holotype lost, original type locality: Paraguay; neotype MNHN Paraguay 6608, type locality: Colonia Ybycui, Estancia Ybycui, Departamento Canindeyú, Paraguay, according to designation by Motte and Cacciali, 2009); Vanzolini and Williams, 1970:8; Peters and Donoso-Barros, 1970:60; Motte and Cacciali, 2009:19; Langstroth, 2006:154; Nicholson et al. 2006:2.

Anolis holotropis Boulenger, 1895:522 (holotype unknown, type locality: "Province Matto Grosso, Brazil").

Norops sladeniae Boulenger, 1903:69 (holotype unknown, type locality: "Chapadá, Matto Grosso" = Chapada dos Guimarães, Mato Grosso, Brasil, according to Vanzolini and Williams, 1970).

Anolis steinbachi Griffin, 1917:308 (holotype CM 988, type locality "Provincia del Sara, Bolívia").

Norops marmorata Amaral, 1932:63 (holotype MZUSP 737, type locality "Jaguará, Rio Grande, Minas Gerais, Brasil").

Anolis chrysolepis meridionalis; Hellmich, 1960:22 (partim, according to Vanzolini and Williams, 1970).

Norops meridionalis; Nicholson 2002:97.

Abbreviated Description. Maximum SVL 56 mm. Specimens analyzed presented vertebral scales anteriorly small, keeled, posteriorly increasing in size and forming a gradual transition toward the granules on flanks; they do not form rows of enlarged scales along dorsum. The neotype description provided by Motte and Cacciali (2009) states that vertebral scales form eight rows of enlarged scales along back that are distinctly larger than scales on the flanks. Scales on upper arm keeled, markedly larger than

vertebral scales. Scales on snout anteriorly weakly to distinctly keeled, uni- or multicarinate and slightly larger than the flat posterior scales. Supraorbital semicircles with enlarged, weakly keeled scales. Supraocular region with a group of distinctly enlarged scales surrounded posteriorly and laterally by granules and anteriorly by smaller scales. Interparietal scale distinctly larger than adjacent scales. Loreals 6–7; canthals 5–6; 8–10 scales between second canthals; 0–1 scales between supraorbital semicircles; 0–2 scales between supraorbital semicircles and interparietal, postmentals 5–8 (Fig. 9F).

Color in Preservative. No sexual dimorphism in color pattern observed. Dorsal color usually grayish-brown, with a light cream, sometimes tending to reddish or light-orange, vertebral band. Dark brown V-shaped lines (apex directed posteriorly) may be present, as well as dark-brown irregular figures and spots across the limbs and in paravertebral region. A pair of subtriangular dark spots on sacral region frequently present.

Dewlap Color in Life and Preservative. When extended, dewlap skin in preservative blue or grayish-blue in center and pale or cream along rim, with light or dark scales. When not extended, grayish-blue laterally, with the center cream or beige. In life, dewlap skin deep blue in center and orange to pale yellow along rim. Scales may be darker on the center, tending to cream or beige on the border, changing to orange or pale yellow on anterior base of dewlap, along rim (Fig. 8F). Langstroth (2006) presented a dewlap photo of a male *A. meridionalis* from near the Zapocós Reservoir in Bolivia showing a deep blue skin with irregular grayish lines and light or dark scales, tending to grayish-green along rim.

Comparisons with the Other Species from the A. chrysolepis Species Group. *Anolis meridionalis* is sympatric with *A. brasiliensis*, of which it can be distinguished mainly by the digital dilatations on phalanx ii and iii, which are continuous with scales under phalanx i and do not form the prominent border observed in *A. brasiliensis*. *Anolis meridionalis* can also be distinguished from

A. brasiliensis by smaller body size (*A. meridionalis* reaches 56 mm, whereas *A. brasiliensis* reaches 69 mm), by the smaller number of scales between supraorbital semicircles (0–1 in *A. meridionalis* and 0–3 in *A. brasiliensis*), by the smaller number of scales between supraorbital semicircles and interparietal (1–4 in *A. brasiliensis* and 0–2 in *A. meridionalis*), and by the smaller number of fourth finger and toe lamellae (15–20 and 25–32, respectively, in *A. brasiliensis*, 10–12 and 18–24 in *A. meridionalis*). Besides these morphological differences, these species do not occur in the same habitat: *A. meridionalis* is commonly found in open areas densely covered by grass in Brazilian Cerrado (Vanzolini and Williams, 1970), whereas *A. brasiliensis* is a typical inhabitant of gallery forests in the same biome (Vitt et al., 2008). Vitt et al. (2008: 146) found “only two specimens of *A. brasiliensis* outside of forested habitat in typical Cerrado, and both were inside termite nests and inactive.”

Distribution. Central Brazil, Paraguay, and Bolivia.

IDENTIFICATION KEY TO THE SPECIES OF THE *ANOLIS CHRYSOLEPIS* GROUP

- 1a. Digital dilatations under phalanx ii and iii are continuous with scales under phalanx i, not forming a prominent border; dewlap skin blue or grayish-blue in center and orange to yellowish-orange (it can be pale or cream in preserved specimens) along rim, with light or dark scales *A. meridionalis*
- 1b. Digital dilatations under phalanx ii form a prominent border over scales under phalanx I 2
- 2a. A weakly to distinctly double row of enlarged vertebral scales from nape to base of tail; scales on upper arm larger than vertebral scales 3

- 2b. Vertebral area with slightly or distinctly enlarged scales, of which the two vertebral rows may be larger than adjacent ones; scales on upper arm smaller than, to sub equal to, vertebral scales 5
- 3a. A double row of weakly enlarged vertebral scales; scales on upper arm slightly larger than vertebral scales; dewlap red along rim (it can be pale in some specimens), with red or blackish scales, and blue with light scales in the center *A. scypheus*
- 3b. A double row of distinctly enlarged vertebral scales from nape to base of tail; scales on upper arm distinctly larger than vertebral scales 4
- 4a. Supraorbital semicircles with enlarged, keeled scales, forming a pronounced ridge in some specimens; supraocular scales with a group of enlarged scales, surrounded in all their extension by distinctly smaller scales; dewlap skin red, but in preserved specimens it usually appear as pale, with light scales *A. planiceps*
- 4b. Supraorbital semicircles with enlarged and generally smooth scales; most supraocular scales large, grading laterally into smaller scales; dewlap skin in preserved specimens blue or grayish-blue with light or grayish scales *A. brasiliensis*
- 5a. Vertebral area with scales not or only slightly enlarged; interparietal scale distinctly larger than adjacent scales; dewlap skin deep blue or blackish-blue with light and grayish scales *A. bimbiceps*
- 5b. Vertebral area with scales slightly to distinctly enlarged; interparietal subequal to or slightly larger than adjacent scales 6
- 6a. Vertebral area with scales distinctly enlarged along back, numbers of enlarged rows increasing posteriorly; interparietal subequal to or slightly larger than adjacent scales; male dewlap royal blue, dark blue, or blackish, with light scales or blue scales toward the rim; female dewlap skin pale yellow to orange (it can be pale or cream in preserved specimens), and toward the rim scales may be darker *A. chrysolepis*
- 6b. Vertebral area with scales slightly enlarged along back, numbers of enlarged rows increasing posteriorly; interparietal larger than adjacent scales; male dewlap royal blue or blackish-blue, with light scales; female dewlap with a central blue spot, surrounded by a cream area; scales usually light *A. tandai*

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APPENDIX 1

Specimens examined (number of specimens of each species in parentheses).

Anolis chrysolepis (50). **SURINAM:** Sipaliwini: Tafelberg Nature Reserve (MCZ R-154861), **Nassau:** Nassau Plateau (MZUSP 29658), **Saramacca:** 1 km from Coppename River (MCZ R-155217). **FRENCH GUIANA:** Sophie trail S from St. Elie gold pits (MCZ R-77548), lower Matarony River, tributary of Approague River (MCZ R-118644, R-118641, R-110388), Saul (MCZ R-146767), **Sinnamary,** Petit Sant River (MPEG 15829, 19845). **GUIANA:** Onora (MCZ R-63350), Shudikar-won (MCZ R-65349). **BRAZIL:** **Amapá:** **Porto Platon** (MCZ 85010), **Serra do Navio** (MCZ R-79146, R-85009; MPEG 19676, 19678, 19679, 19680, 19681, 19682, 19683, 19685, 19686, 19687, 19688, 19690), **Mazagão:** Rio Maracá (MZUSP 8845, 8846; CHUNB 56761), **Laranjal do Jari** (CHUNB 56767; MZUSP 83222, 83224), **Pará:** Aldeia dos índios Tirió (MZUSP 13141), **Oriximiná:** FLONA Saracá-Taquera: Platô Aviso (MPEG 27114, 27217), Platô Almeidas (MPEG 26864), Estação Ecológica Grão-Pará, Serra do Acari (MPEG 26573, 27374, 27375, 27376, 27377, 26587, 26578, 26593), **Porto Trombetas:** Platô Bela Cruz (MPEG 24216), Platô Greig (MPEG 24218), **Faro:** Floresta Estadual de Faro (MPEG 26597).

Anolis planiceps (72). **VENEZUELA:** **Portuguesa** (MCZ R-176492, R-176491, R-176493, R-176494, R-176495, R-176496): Acarigua: Environs of Agua Blanca (MCZ R-123708), **Distrito Federal:** Puerto la Cruz (MCZ R-48782), **Amazonas:** Puerto Ayacucho (MCZ R-58328, R-84073), **Sucre:** Cunamacoa, San Rafael (MCZ R-48781), Peninsula of Paria, Yacua (MCZ R-43856, R-43860, R-43857, R-43858), **Aragua:** Palsamencho Parque, HPittier between Portochuelo and Ocumare (MCZ R-101817), **Carabobo:** Hacienda San Esteban, North of Puerto Cabello (MCZ R-107688), Goaguaza: Miquija (MCZ R-121759, R-121758, R-121762, R-121760, R-121757), **Salon:** Las Quiguas (MCZ R-121753, R-81302), **Falcón:** Acosta, Pauji, (MCZ R-49035, R-48723, R-49037, R-48725,

R-48724, R-49036), Riecito (MCZ R-49050), **Miranda:** Guatapo (MCZ R-100438), **Bolívar:** Arabupu, Mt. Roraima (MCZ R-34868). **TRINIDAD:** Nariwa Swamp, Cascadoux Trace (MCZ R-60800), Huevas Id. (MCZ R-100119), **Sangre Grande:** Toco (MCZ R-10746, R-10747), La Seiva (MCZ R-8998, R-8999, R-9000), **Trinidad:** Chacachacare (MCZ R-100118), Maruga (MCZ R-31495), Palmiste: 3 km S San Fernando (MCZ R-85011, R-85012, R-85013, R-85014, R-85015, R-85016, R-85017, R-85018, R-100120, R-100121, R-100122, R-81303, R-81304, R-81501, R-85474, R-85475, R-85473). **BRAZIL:** **Amazonas:** Serra da Neblina (MCZ R-86763), Manaus: Reserva Ducke (MCZ R-92682), Camp Gavião, approx. 100 km N of Manaus (MCZ R-168987). **GUYANA:** **Pomeroon-Supenaam:** Dawa (MCZ R-123745), **Cuyuni-Mazaruni:** Kaburi, 30 miles from Bartica (MCZ R-81306), Mazaruni River (MCZ R-39690), Kamakusa (MCZ R-65352).

Anolis scypheus (49). **PERU:** **Loreto:** (KU 222147), Rio Pacaya, Cahuana (MCZ R-160775), Galicia: west bank of Rio Tapiche (MCZ R-157245), Pampa Hermosa: near mouth Cushcubatey River (MCZ R-57369, R-57373, R-57372), **Amazonas:** Mouth of R.Santiago Marañon (MCZ R-57371), **Ucayali:** River Tamaya (MCZ R-57374), **Madre de Dios:** Cocha Cashu, NW of mouth of Rio Manu (MCZ R-178177). **ECUADOR:** **Napo:** Trail from Laguna Taracoa, S bank Rio Napo, 30 km downriver from Coca (MCZ R-154571), Hacienda 'Primavera' N bank Rio Napo 30 km from Coca (MCZ 92612, 154567, 154568), Prov. S side of Rio Napo, 6.5 km ESE of Puerto Misahualli, at La Cruz Blanca on ASuarez's land (MCZ R-171889), **Sucumbios:** Santa Cecilia (MCZ R-92601, R-92602, R-92603, R-92604, R-92605, R-92608), Santa Cecilia, Rio Agvaratis (MCZ R-100010), Limon Cocha (MCZ R-156822, R-85098, R-85099, R-85100, R-85101, R-92610, R-92611, R-92612), **Chimborazo:** Riobamba (MCZ R-29290). **COLOMBIA:** **Meta:** Villavicencio: San Martin (MCZ R-32309, R-32310, R-32312, R-32313, R-32314, R-32315, R-32317), Bairro Povenir (MZUSP 36099, 36100). **BRAZIL:** **Amazonas:** Acanauí (MZUSP 47212, 47214), Rio Japurá: Serrinha (MZUSP 46793, 46794), Costa da Altamira (MZUSP 47301, 47302, 47303); Lago Amanã (MZUSP 60462).

Anolis tandai (96). **BRAZIL:** **Rondônia:** Espigão do Oeste (MPEG 21479, 21483, 21488, 21899), **Amazonas:** Estirão do Equador (MPEG 901), Rio Urucu (holotype: MPEG 15850), Benjamin Constant (paratypes: MPEG 15933, 15938, 15949, 15986, 15987, 15995; MZUSP 15898), Madeireira Sheffer, Rio Ituxi (MPEG 20372, 20473, 20474), Careiro da Várzea, estrada para Altazes (MPEG 18918, 18920, 18928, 18931, 18934, 18935), Tapauá (MZUSP 37763, 37764, 37765), Berurú (MZUSP 38101, 38102), Novo Aripuanã (MZUSP 42400), Borba (MZUSP 41077, 9145), Rio Jurua (MZUSP 700), Barreira do Matupiré, Rio Madeira, (MZUSP 42150), Maués, Bragança/São Tomé, Rio Paraconi (MPEG 27673, 27674, 27675, 27676), **Acre:** Rio Jurua, approx. 30 km North of Porto Walter of Walter (MPEG 20655, 20656, 20658, 20659,

20662, 20663, 20665, MZUSP 53270, 53271, 53272), Alto Purus (MZUSP 2413, 2514), Seringal Santo Antônio, próximo a Manoel Urbano (MZUSP 32097), Parque Nacional Serra do Divisor, Estirão do Panela (MZUSP 88656), Estirão do Equador (MZUSP 899), **Pará:** Parque Nacional da Amazônia, Itaituba (MPEG 21986, 21987, 21988, 21989, 21990, 21991), Parque Nacional da Amazônia, Uruá (MZUSP 52533, 52536, 52538, 52539), Cachoeira da Montanha (MZUSP 53657, 53658), Buburé (MZUSP 53692, 53693, 53695), Barreirinha, próximo a São Luís, Rio Tapajós (MZUSP 20679), **Mato Grosso:** Aripuanã (MZUSP 82590, 82585, 82589, 82587, 82591, 82593, 82586, 82592, 82588, 82594, 81502, 81512, 81501, 81520, 81505, 81497, 81508, 81514, 81491, 81491), Juruena (MZUSP 82402, 82404, 82397, 82403, 82400). **PERU:** **Loreto:** Rio Orosa (MZUSP 56658), Alto Curanja, Igarapé Champuia (MZUSP 3324, 3324).

Anolis brasiliensis (116). **BRAZIL:** **Mato Grosso:** Barra do Tapirapés (holotype: MZUSP 10319, paratypes: MCZ R-98284, R-98285, R-98286, R-98287, R-98288, R-98289), Porto Alegre do Norte (CHUNB 47842), Vila Rica (MZUSP 82874, 82884, 82882), **Tocantins:** Caseara (CHUNB 44989, 44992, 44499, 45000, 45009, 45036, 45028, 45026, 45022, 45030), Pium (CHUNB 24754, 24756), Palmas (CHUNB 25082, 24216, 11521, 11522, 24215, 16947, 25092, 11520, 24262, 15230, 16142, 16948), Peixe (CHUNB 52471, 52472, 52474), Porto Alegre do Tocantins (CHUNB 38918), Porto Nacional (CHUNB 47741, 47742, 47740), Mateiros (CHUNB 27161, 27158, 27162), Usina Hidroelétrica Luis Eduardo Magalhães (MZUSP 95466, 95467), **Goiás:** São Domingos (CHUNB 35282, 35257, 35288, 35319, 35285, 35314, 43823, 35315, 35311, 35258, 43824, 43832, 33020), Minaçu (CHUNB 11019, 11021, 10967, 10983, 49706, 48617, 10987, 49703, 10986, 10970, 10999, 08655, 29309), Colinas do Sul (CHUNB 50396, 50393, 50395, 44694), Valparaíso (CHUNB 08556, 08555), Luziânia (CHUNB 47427, 43408, 40865, 43407, 47426), Três Ranchos (CHUNB 44740), Goiânia (CHUNB 57318), **Minas Gerais:** Unai (CHUNB 30888, 32873, 32867, 36287), **Distrito Federal:** Brasília (CHUNB 49614), **Piauí:** Ribeiro Gonçalves (CHUNB 57028), **Maranhão:** Carolina (CHUNB 51972, 51973, 51974, 51975), Gancho do Arari (MZUSP 60682), **Pará:** Novo Progresso (CHUNB 34542), Parauapebas (CHUNB 08855, 11132), Canaã dos Carajás (MPEG 25166), Itaituba, Jacareacanga (CHUNB 56398), **Ceará:** Arajara (MZUSP 51698, 51699, 51703, 51706), **São Paulo:** Bueno de Andrade (MZUSP 4384), Ibarra (MZUSP 4487), Itapura (MZUSP 551).

Anolis meridionalis (9). **BRAZIL:** **Minas Gerais:** Paracatu (CHUNB 26140, 26146), Buritizeiro (CHUNB 44522), **Rondônia:** Vilhena (CHUNB 11715), **Tocantins:** Palmas (CHUNB 12022, 12026), **Goiás:** Luziânia (CHUNB 43404, 43405), **Distrito Federal:** Brasília (CHUNB 43282).

Anolis bimbiceps (11). **PERU:** **Loreto** (MZUSP 13135, 131137, 13138), N.A. 1.5 km N of Teniente Lopez, 310 (KU 222145). **COLOMBIA:** **Amazonas:**

6 km NW of Leticia (MCZ R-112267). **ECUADOR:** **Pastaza:** Conambo (MZUSP 13140). **BRAZIL:** **Amazonas:** Apuí (CHUNB 38340); São Gabriel da Cachoeira: Missão Taraquá, Rio Uapés (MPEG 17819), Uatumã (MZUSP 17439), Igarapé Belém, Rio Solimões (MZUSP R-13026, 13027).

Anolis onca (1). **VENEZUELA:** **Falcón:** Rio Seco on Caribbean Sea, between Urumaco and Coro (MCZ R-132736).

Anolis lineatus (1). **NETHERLANDS ANTILLES:** **Curaçao:** Cos Cora (MCZ R-83027).

Anolis sagrei sagrei (1). **JAMAICA:** **Westmoreland:** Savanna La Mar (MCZ R-161754).

Anolis auratus (1). **COLOMBIA:** **Magdalena:** Pozo Colorado, 11 km W of Santa Maria (MCZ R-104546).

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