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# A Phylogenetic Analysis of *Vanzolinius* Heyer, 1974 (Amphibia, Anura, Leptodactylidae): Taxonomic and Life History Implications

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## A PHYLOGENETIC ANALYSIS OF *VANZOLINIUS* HEYER, 1974 (AMPHIBIA, ANURA, LEPTODACTYLIDAE): TAXONOMIC AND LIFE HISTORY IMPLICATIONS <sup>1</sup>

(With 1 figure)

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**ABSTRACT:** The validity of the monotypic leptodactylid frog genus *Vanzolinius* Heyer, 1974 has been questioned recently. We explore the relationships of *Vanzolinius discodactylus* within the cluster of closely related genera *Adenomera*, *Leptodactylus*, and *Lithodytes* with both morphological and molecular data sets. Morphological and combined morphological and molecular data were analyzed using maximum parsimony; molecular data sets were analyzed with maximum likelihood methods. The resultant relationships are unambiguous in *Vanzolinius* being imbedded within *Leptodactylus*. In order to maintain *Leptodactylus* as a monophyletic genus, *Vanzolinius* is placed in the synonymy of *Leptodactylus* Fitzinger, 1826. The implications of relationships analyzed in this study are discussed in terms of both nomenclature and life-history evolution.

**Key words:** *Leptodactylus*. *Vanzolinius*. Phylogenetic relationships. Life history evolution.

**RESUMO:** Análise filogenética de *Vanzolinius* Heyer, 1974 (Amphibia, Anura, Leptodactylidae): implicações taxonômicas e sobre a história de vida.

A validade do gênero monotípico de leptodactídeo *Vanzolinius* Heyer, 1974, tem sido questionada recentemente. Neste estudo exploramos as relações de *Vanzolinius discodactylus* dentro do agrupamento de gêneros proximamente relacionados *Adenomera*, *Leptodactylus* e *Lithodytes* por meio de dados morfológicos e moleculares. Dados morfológicos e dados morfológicos e moleculares combinados foram analizados por parcimônia máxima, dados moleculares foram analisados por máxima verossimilhança. As relações resultantes são inequívocas em *Vanzolinius* ter que ser incluído em *Leptodactylus*. Para manter *Leptodactylus* como um gênero monofilético, *Vanzolinius* Heyer 1974, é colocado na sinonímia de *Leptodactylus* Fitzinger, 1826. As implicações dos relacionamentos analisados neste estudo são discutidas em termos de nomenclatura e evolução dos modos reprodutivos.

**Palavras-chave:** *Leptodactylus*. *Vanzolinius*. Relações filogenéticas. Evolução da história de vida.

### INTRODUCTION

The frog genera *Adenomera* Fitzinger, 1867, *Lithodytes* Fitzinger, 1843, and *Vanzolinius* Heyer, 1974 have, at one time or another, been included in the genus *Leptodactylus*. BOULENGER (1883) described the currently recognized monotypic *Vanzolinius* as *Leptodactylus discodactylus*. HEYER (1970) associated this taxon with the *Leptodactylus melanotus* species group. Later, HEYER (1974a) placed the taxon within *Lithodytes* commenting on its possible distinctiveness and subsequently created the genus *Vanzolinius* to accommodate this species

(HEYER, 1974b). The most recent morphological analysis indicated that *Vanzolinius* shared distinctive characteristics with *Leptodactylus diedrus* (HEYER, 1998). Previous analyses of relationships agreed that within the subfamily Leptodactylinae the genera *Adenomera*, *Leptodactylus*, *Lithodytes*, and *Vanzolinius* formed a monophyletic clade and that the genus *Physalaemus* Fitzinger, 1826, was more distantly related to this clade (HEYER, 1974a, 1975; LYNCH, 1971).

It is necessary to establish convincingly whether the genus *Leptodactylus* as currently understood is monophyletic, if we wish to understand the

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evolution of life history variation in *Leptodactylus*. In this paper, we are particularly interested in determining the phylogenetic relationships of *Vanzolinius*. Preliminary findings on relationships of previously proposed monophyletic clades in the *Leptodactylus* cluster (*Adenomera*, *Leptodactylus*, *Lithodytes*, *Vanzolinius*) are also presented in this paper, and we discuss the implications of our results for understanding aspects of life history evolution in this cluster.

#### MATERIAL AND METHODS

**Taxon sampling** – Species groups within *Leptodactylus* were previously recognized on the basis of morphological and life history characters (HEYER, 1969). We included samples from each of the four species groups to sample the morphological diversity within *Leptodactylus*. *Leptodactylus riveroi* Heyer & Pyburn, 1983, a species of uncertain species group affinity, and *L. silvanimbus* McCranie *et al.*, 1980, a species recently suggested as basal within the genus (HEYER, DE SÁ & MULLER, 2005), were also included. *Physalaemus* has been shown to function well as an outgroup for *Leptodactylus* using both morphological and molecular data (HEYER, 1998; HEYER, DE SÁ & MULLER, 2005); herein *Physalaemus gracilis* (Boulenger, 1883) was the outgroup taxon.

The taxa analyzed in this study are: *Leptodactylus bufonius* Boulenger, 1894, *L. fuscus* (Schneider, 1799) (*fuscus* species group); *L. leptodactyloides* (Andersson, 1945), *L. melanotus* (Hallowell, 1861) (*melanotus* species group); *L. chaquensis* Cei, 1950, *L. insularum* Barbour, 1906 (*ocellatus* species group); *L. pentadactylus* (Laurenti, 1768) (*pentadactylus* species group); *L. diedrus* Heyer, 1994, *L. riveroi*, *L. silvanimbus* (*Leptodactylus* of unclear species group affinity); *Adenomera hylaedactyla* (Cope, 1868), *Lithodytes lineatus* (Schneider, 1799), *Vanzolinius discodactylus* (Boulenger, 1883); and *Physalaemus gracilis* (as the outgroup). For both the morphological and molecular data, the data for *L. pentadactylus* are from Middle American specimens. See Tissue Voucher Specimens section at the end of this paper for specimen data used for molecular analyses. Museum abbreviations follow LEVITON *et al.* (1985).

**Morphological data set** – The morphological matrix is provided in Appendix 1. The character state descriptions and ordering information are the same as those published in HEYER (1998) with the

following exceptions. We had no tissue samples for *Adenomera marmorata* and *Physalaemus pustulosus*, two of the taxa used in HEYER (1998), so we used morphological data for *Adenomera hylaedactyla* and *Physalaemus gracilis*, for which we do have molecular data. Data taken for *A. hylaedactyla* and *P. gracilis* were taken from HEYER (1974a), HEYER, DE SÁ & MULLER (2005), USNM 292477 (cleared-and-stained *A. hylaedactyla*) and RdS 511 (larval *P. gracilis* from Uruguay, Canelones, Balneario Atlantida, Rafael de Sá field number). These two species have a few states that differ from their congeners, and require recoding of states and/or redefinition of states as follows.

Character 7, toe webbing. *Physalaemus pustulosus* was coded as having a unique state in the data set of HEYER (1998), toes with weak basal fringes and webbing. *Physalaemus gracilis* has toes without web or fringes, a condition found in other taxa in the data set. The new definitions are: State 0 – toes without web or fringes; State 1 – toes with fringes extending length of toes except for tips; State 2 – females with weakly developed lateral toe fringes and males either with ridges or weakly developed fringes. The state ordering is 0-1-2.

Character 15, depressor mandibulae muscles. The depressor mandibulae may have one to three slips of origin, from the dorsal fascia (df), the zygomatic and/or otic ramus of the squamosal (sq), and the tympanic annulus (at) (following the terminology defined by STARRETT, 1968). Lower case indicates small slips of the muscle, upper case indicates large slips. *Physalaemus pustulosus* has the dfSQat condition, whereas *P. gracilis* has DFSQat. The DFSQat condition is state 0 in our data matrix.

Character 18, anterior petrohyoideus muscle. *Adenomera hylaedactyla* has a state not found in the data set of HEYER (1998). The new definitions are: State 0 – the anterior petrohyoideus muscle inserts entirely on the edge of the hyoid apparatus; State 1 – the muscle inserts on the edge of the hyoid and on the ventral body of the hyoid in part; State 2 – the muscle inserts entirely on the ventral surface of the hyoid body. The state ordering is 0-1-2.

Character 24, sartorius muscle. The condition in *P. gracilis* does not differ from some other taxa in the data set, in contrast to the condition found in *P. pustulosus*. The new definitions are: State 0 – muscle moderate; State 1 – intermediate condition between States 0 and 2; State 2 – muscle broad. The state ordering is 0-1-2.

Character 32, sacral diapophyses. *Physalaemus gracilis* does not differ in this character from other taxa. Thus characters 32-37 in our data set equal characters 33-38 in the HEYER (1998) data set. Molecular methodology – DNA extraction followed HILLIS *et al.* (1996). Two segments of the mitochondrial genome were amplified using the polymerase chain reaction (PCR). A segment of the 12S r RNA of ~ 900 nucleotides and a segment of the 16s r RNA of ~ 700 nucleotides were amplified. Double-stranded (DS) PCR amplifications were performed in a final volume of 50 $\mu$ l containing 0.4 $\mu$ l of each primer, 1.0 $\mu$ l of each dNTP, 3.0 $\mu$ l of 25mM MgCl, and 1.25 units of *Taq* (*Thermus aquaticus*) DNA polymerase; the reaction was overlaid with 50 $\mu$ l of mineral oil. PCR conditions were as follows: 94°C for 60s, 57°C for 60s, and 72°C for 60s, with 25 cycles for the 12S amplification and 30 cycles for the 16S amplification. Amplified product was purified using Wizard® PCR Preps Kit (Promega). Of the purified DS fragment, 0.5 $\mu$ l were mixed with 1.5 $\mu$ l of a single IRD-labeled primer, 7.2 $\mu$ l of Sequencing Buffer, 1.0 $\mu$ l of Sequitherm Excel™II (Epicentre Technologies Co.) DNA polymerase, and 6.8 $\mu$ l of dH<sub>2</sub>O. Subsequently, 4.0 $\mu$ l of this mix was added to each of 4 tubes containing 2 $\mu$ l of each nucleotide respectively. PCR conditions were as follows (30 cycles): 92°C for 30s, 55°C for 30s, and 70°C for 30s. SS amplified and IR labeled fragments were sequenced in a LI-COR 4200 IR DNA Sequencer on 6% acrylamide gels. A total of 839 12S and 648 16S nucleotide positions were aligned unambiguously using Clustal X and positions of ambiguous alignments were not used in the phylogenetic analyses. GenBank accession numbers for the sequence data are AY943217–242. The alignment matrix is provided in Appendix 2.

Phylogenetic Analysis – Maximum Parsimony (MP) analysis using PAUP\* 4.0 (SWOFFORD, 2002) was used for both the morphological data set and the combined morphological and molecular data set. Molecular data sets were analyzed with maximum likelihood (ML) in PAUP\* under the GTR+I+G model recommended by both the Hierarchical Likelihood Ratio Test and the Akaike Information Criterion used by Modeltest 3.04 (POSADA & CRANDALL, 1998). We obtained a total of 37 morphological characters and 1486 base pairs (bp) for each taxon (839 bp corresponding to the 12S rDNA gene and 647 bp to the 16S rDNA gene). Sequences were aligned using Clustal X (THOMPSON, HIGGINS & GIBSON, 1994). We ran individual analyses for each of the

data sets (i.e., morphology, 12S, and 16S data sets) as well as combined analyses (i.e., 12S+16S matrix, morphology+12S+16S matrix). In combined analyses gaps were alternatively considered as missing or as 5th characters; we also evaluated the effect of the substitution bias in the analysis of the combined data matrix using MP by down-weighting transitions to transversions 5:1.

## RESULTS

There is modest variation in the 12S, 16S, and 12S+16S data sets (Tabs.1-3). The maximum sequence divergences between pairs of taxa are 21% for the 12S data, 16% for the 16S data, and 18% for the 12S+16S data.

The results of all cladistic analyses are almost identical; consequently we present the maximum parsimony combined data set results and point out where the analyses differ (Fig.1). The parsimony analysis of the combined data matrix results in a single tree (length=1430, consistency index=0.56) in which *Vanzolinius* exhibits a sister taxa relationship with *L. diedrus*. This relationship is also recovered in the analyses of the combined molecular data partitions as well as in all analyses of the 12S data partition. The analyses of the 16S data partition position *Vanzolinius* in the following clade (*L. diedrus* (*L. leptodactyloides*+*Vanzolinius*)). The distance data matrices show that the close relationship of *L. diedrus* with *Vanzolinius* is unambiguous in the 12S data (Tab.1), but not at all clear in the 16S data, where *L. diedrus* and *Vanzolinius* have lower sequence distance values with *L. silvanimbus* and several members of the *L. fuscus*, *L. melanotus*, and *L. ocellatus* group members than with each other (Tab.2). The morphological data set demonstrates strong support for a *L. diedrus*-*V. discodactylus* sister species relationship with 100% bootstrap support.

## DISCUSSION

Phylogenetic conclusions – The following conclusions are supported by the analyses performed on our data.

First, *Vanzolinius* always clusters within *Leptodactylus*. The data are very clear and convincing for this conclusion. There are two nomenclatural options to resolve the phylogenetic conclusion that *Vanzolinius* is imbedded within *Leptodactylus*: *Vanzolinius* could be synonymized with *Leptodactylus*; or one or more clades within

*Leptodactylus* could be raised to generic status. Current (unpublished) data are inconclusive regarding the phylogenetic relationships among *Leptodactylus* species, and rule out elevating certain clades within *Leptodactylus* to generic status at this time. However, we think there are compelling arguments for placing *Vanzolinius* in the synonymy of *Leptodactylus*. The previous actions on generic placement of the species *discodactylus* were all based on morphological and karyotype data. The strongest support for generic recognition of *Vanzolinius* as a genus distinct from *Leptodactylus* involved two morphological features of the toes: the toe tips of *V. discodactylus* are expanded into small disks with longitudinal grooves on the dorsal surface and the terminal phalanges are T-shaped (HEYER, 1974b). With the discovery of *Leptodactylus diedrus*, the morphological distinctiveness between *Leptodactylus* and *Vanzolinius* was bridged to a large extent (HEYER, 1998). Thus, the morphological data used to define *Vanzolinius* as a genus distinct from *Leptodactylus* are seriously compromised by inclusion of the data for *L. diedrus* and the molecular data strongly support synonymizing *Vanzolinius* with *Leptodactylus*. Consequently, we hereby synonymize the genus *Vanzolinius* Heyer, 1974 with the genus *Leptodactylus* Fitzinger, 1826.

Second, the genera *Adenomera* and *Lithodytes* may share a sister-group relationship and our data provide support that both are evolutionarily distinct from *Leptodactylus* (including *Vanzolinius*).

Third, the previously recognized “traditional” species groups may not all be monophyletic, although the two members of the *L. fuscus* group form a well-supported clade in this study.

Fourth, a sister-group relationship between *L. discodactylus* and *L. diedrus*, previously suggested by HEYER (1998), is reasonably well supported by the morphological and combined molecular data sets.

Finally, *Leptodactylus riveroi*, a taxon of uncertain relationships, exhibits suggestive affinities to the *L. melanotus* species group.

**Life history implications –** All members of the subfamily Leptodactylinae (except *Limnomedusa*), place their eggs in foam nests (LANGONE, 1995). Within the *Leptodactylus* cluster, however, there is variation regarding where the foam nests are deposited and considerable variation occurs in other life history aspects. Two examples illustrate how an understanding of phylogenetic relationships in this group is critical to deciphering life history evolution in the genus *Leptodactylus*.

Table 1. 12S sequence differences between taxon pairs included in study using General Time Reversible (GTR) parameter values.

TAXA	1	2	3	4	5	6	7	8	9	10	11	12	13	14
1 <i>L. diedrus</i>	-													
2 <i>L. riveroi</i>	0.148	-												
3 <i>L. silvaniimbis</i>	0.126	0.137	-											
4 <i>L. bufonius</i>	0.139	0.144	0.115	-										
5 <i>L. fuscus</i>	0.136	0.165	0.135	0.077	-									
6 <i>L. chaguenensis</i>	0.128	0.119	0.078	0.095	0.097	-								
7 <i>L. insulatum</i>	0.123	0.133	0.094	0.094	0.096	0.065	-							
8 <i>L. leptodactyloides</i>	0.131	0.135	0.086	0.107	0.116	0.042	0.087	-						
9 <i>L. melanotus</i>	0.137	0.146	0.101	0.105	0.117	0.087	0.088	0.097	-					
10 <i>L. pentadactylus</i>	0.144	0.160	0.116	0.118	0.118	0.107	0.113	0.115	0.131	-				
11 <i>V. discodactylus</i>	0.113	0.166	0.141	0.136	0.129	0.126	0.116	0.130	0.136	0.134	-			
12 <i>A. hylaedactyla</i>	0.177	0.197	0.177	0.156	0.157	0.145	0.151	0.156	0.174	0.161	0.168	-		
13 <i>Lith. lineatus</i>	0.207	0.203	0.175	0.168	0.173	0.175	0.187	0.178	0.182	0.165	0.190	-		
14 <i>P. gracilis</i>	0.185	0.212	0.167	0.151	0.156	0.161	0.164	0.162	0.171	0.182	0.160	0.174	-	

Table 2. 16S sequence differences between taxon pairs included in study using General Time Reversible (GTR) parameter values.

TAXA	1	2	3	4	5	6	7	8	9	10	11	12	13	14
1 <i>L. diedrus</i>	-													
2 <i>L. riveroi</i>	0.129	-												
3 <i>L. silvanimbus</i>	0.099	0.111	-											
4 <i>L. bufonius</i>	0.118	0.136	0.101	-										
5 <i>L. fuscus</i>	0.096	0.132	0.099	0.053	-									
6 <i>L. chaquensis</i>	0.085	0.105	0.072	0.093	0.079	-								
7 <i>L. insularum</i>	0.088	0.098	0.060	0.088	0.072	0.039	-							
8 <i>L. leptodactyloides</i>	0.092	0.124	0.109	0.117	0.116	0.085	0.092	-						
9 <i>L. melanotus</i>	0.083	0.100	0.079	0.097	0.091	0.060	0.063	0.084	-					
10 <i>L. pentadactylus</i>	0.108	0.128	0.099	0.096	0.082	0.086	0.074	0.117	0.080	-				
11 <i>V. discodactylus</i>	0.104	0.118	0.125	0.132	0.114	0.099	0.094	0.087	0.100	0.114	-			
12 <i>A. hylaeactyla</i>	0.135	0.136	0.128	0.144	0.133	0.128	0.124	0.131	0.111	0.128	0.147	-		
13 <i>Lith. lineatus</i>	0.155	0.146	0.116	0.138	0.125	0.120	0.156	0.121	0.131	0.160	0.105	-		
14 <i>P. gracilis</i>	0.160	0.163	0.126	0.150	0.139	0.126	0.124	0.148	0.145	0.143	0.165	0.143	0.133	-

Table 3. Combined 12S &amp; 16S sequence differences between taxon pairs included in study using General Time Reversible (GTR) parameter values.

TAXA	1	2	3	4	5	6	7	8	9	10	11	12	13	14
1 <i>L. diedrus</i>	-													
2 <i>L. riveroi</i>	0.140	-												
3 <i>L. silvanimbus</i>	0.114	0.125	-											
4 <i>L. bufonius</i>	0.130	0.140	0.109	-										
5 <i>L. fuscus</i>	0.118	0.150	0.118	0.066	-									
6 <i>L. chaquensis</i>	0.109	0.113	0.075	0.094	0.089	-								
7 <i>L. insularum</i>	0.108	0.118	0.079	0.092	0.085	0.054	-							
8 <i>L. leptodactyloides</i>	0.114	0.130	0.096	0.111	0.116	0.060	0.089	-						
9 <i>L. melanotus</i>	0.113	0.126	0.092	0.101	0.106	0.075	0.077	0.091	-					
10 <i>L. pentadactylus</i>	0.128	0.146	0.109	0.108	0.102	0.098	0.096	0.116	0.108	-				
11 <i>V. discodactylus</i>	0.109	0.144	0.134	0.123	0.114	0.106	0.111	0.120	0.125	-				
12 <i>A. hylaeactyla</i>	0.158	0.170	0.155	0.151	0.147	0.138	0.139	0.145	0.146	0.146	0.159	-		
13 <i>Lith. lineatus</i>	0.184	0.178	0.148	0.155	0.152	0.153	0.157	0.168	0.154	0.150	0.177	0.136	-	
14 <i>P. gracilis</i>	0.174	0.190	0.149	0.150	0.149	0.145	0.147	0.156	0.155	0.160	0.175	0.152	0.156	-

First, two clades (*Adenomera* and the *L. fuscus* species group) within Leptodactylinae share the same pattern of males constructing a terrestrial subsurface chamber, attracting females to the chamber acoustically, and depositing the foam nest in the chamber where at least embryonic and early larval development take place (see KOKUBUM & GIARETTA, 2005 and references cited therein). Our data indicate that this complex life history pattern was independently derived in both clades and is not the result of shared ancestral adaptations. Also, at least some members of the *L. pentadactylus* group use pre-existing terrestrial burrows in which they deposit their foam nest (see GIBSON & BULEY, 2004 and references cited therein). Additional taxon sampling is required to determine whether this pattern served as a precursor to the actual construction of terrestrial incubating chambers in the *L. fuscus* group. Our preliminary data suggest support for this scenario.

Second, there is considerable variation in female attendance of foam nests and larvae, whether attending females communicate with their larvae, and how females communicate with their larvae (VAZ-FERREIRA & GEHRAU, 1975; WELLS & BARD, 1988). As far as is known, parental care does not occur in any species of the *L. fuscus* group. Our preliminary data indicate that intensive taxon sampling with additional data is required to resolve relationships among the *Leptodactylus* species that demonstrate female attendance and communication with their offspring in order to understand the evolution of parental care in *Leptodactylus*.

More intensive taxon sampling and the sequencing of nuclear and more slowly evolving genes should provide a well-supported phylogeny for *Leptodactylus* at the species level that will allow a better understanding of the evolution of life history variation in the *Leptodactylus* cluster.

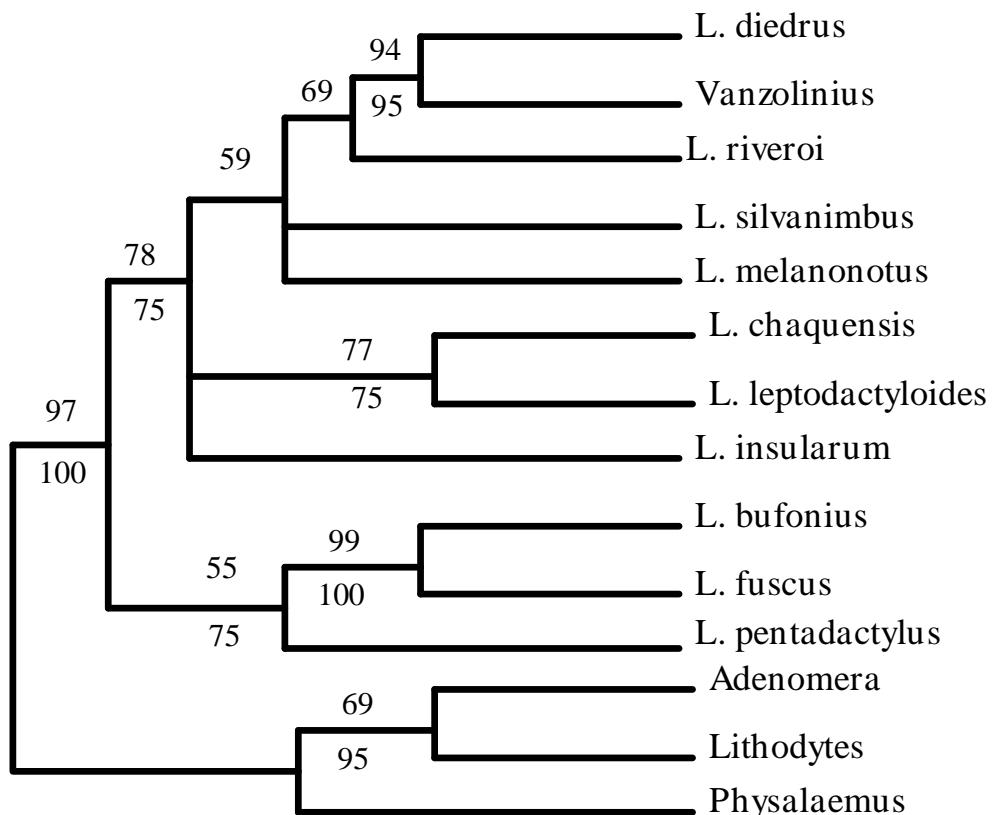


Fig. 1- Maximum Parsimony Tree of combined (morphological and molecular) data sets. Gaps were considered as a fifth character. Numbers above branches correspond to bootstrap support in parsimony analysis; numbers below branches are bootstrap support values from Maximum Likelihood analysis of the combined molecular data set.

## TISSUE VOUCHER SPECIMENS

*Adenomera hylaedactyla* – BRAZIL: PARÁ: Alter do Chão (MZUSP 70958)

*Leptodactylus bufonius* – ARGENTINA: SALTA: 54 km NE of Joaquín V. González on provincial route 41 (USNM field number 175816, deposited in FML).

*Leptodactylus chaquensis* – ARGENTINA: TUCUMÁN: ca 40 km SE San Miguel de Tucumán at km post 1253 on International Route 9 (USNM 319708).

*Leptodactylus diedrus* – VENEZUELA: AMAZONAS: Río Negro, near Neblina base camp on left bank of Río Baria (= Río Mawarinuma) (USNM 30715).

*Leptodactylus discodactylus* – ECUADOR (QCAZ 16788).

*Leptodactylus fuscus* – BRAZIL: RORAIMA: Caracaranã, near Normandia (MZUSP 67073).

*Leptodactylus insularum* – PANAMA: PANAMA: Río Indio, camino hacia Las Minas (CH 4956).

*Leptodactylus leptodactyloides* – BRAZIL: PARÁ: Serra de Kokoinhokren (MZUSP 70969).

*Leptodactylus melanotus* – BELIZE: CAYO: between San Jacinto and Spanish Lookout road on Webster Highway, Caesar's Hotel (USNM 535964).

*Leptodactylus ocellatus* – BRAZIL: SANTA CATARINA: Campeche (MZUSP 68993).

*Leptodactylus "pentadactylus"* – PANAMA: BOCAS DEL TORO: Isla Popa (USNM 347153).

*Leptodactylus riveroi* – VENEZUELA: AMAZONAS: Río Negro, Neblina base camp on left bank of Río Baria (= Río Mawarinuma) (USNM 562029).

*Leptodactylus silvanimbus* – HONDURAS: OCOTEPEQUE; Belén Gualcho (USNM 348631).

*Lithodytes lineatus* – BRAZIL: MATO GROSSO: Apiacás (MZUSP 80874).

*Physalaemus gracilis* – URUGUAY: SALTO: Espinillar (RdS 788 field number).

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## APPENDIX 1

## Morphological (primarily) data matrix used for phylogenetic analysis

Characters	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19
<i>L. bufonius</i>	3	0	0	0	0	0	0	0	3	4	0	3	0	1	1	0	2	0	0
<i>L. fuscus</i>	3	0	0	3	0	0	0	0	3	4	0	3	0	1	1	0	1	0	1
<i>L. leptodactyloides</i>	1	0	2	1	1	0	1	0	3&5	0	2	0	0	0	1	0	0	0	0
<i>L. melanotonus</i>	1	0	2	0	0	0	1	1	5	0	2	0	0	0	1	0	0	0	0
<i>L. chaquensis</i>	2	0	2	3	0	0	1	2	2	0	2	0	0	0	1	0	0	0	0
<i>L. insularum</i>	1	0	2	2	0	0	1	1	3	0	2	0	0	0	1	0	0	0	1
<i>L. pentadactylus</i>	1	0	1	2	0	0	0	2	4	0	3	1	0	0	1	0	0	0	0
<i>L. diedrus</i>	1	0	2	0	2	0	1	?	?	?	?	1	?	1	0	0	0	0	0
<i>L. riveroi</i>	0	0	2	2	0	0	1	0	4	0	2	0	?	?	1	0	0	0	0
<i>L. silvanimimus</i>	1	0	2	0	0	2	0	5	0	2	0	0	?	1	0	0	0	0	0
<i>A. hylaedactyla</i>	1	0	0	1	0	0	?	0	0	1	0	1	0	1	1	1	1	1	0
<i>Lith. lineatus</i>	1	0	0	2	4	0	0	?	1	0	0	0	1	1	1	0	0	0	1
<i>V. discodactylus</i>	1	0	0	3	0	1	0	5	0	2	0	1	0	1	0	0	1	0	1
<i>P. gracilis</i>	4	1	3	0	0	1	0	3	4	1	3	0	1	0	0	1	2	0	0
<i>L. bufonius</i>	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	37
<i>L. fuscus</i>	0	0	1	0	2	0	0	1	2	1	1	0	0	0	0	2	1	1	0
<i>L. leptodactyloides</i>	0	0	1	0	0&2	0	0	0	2	1	2	0	0	0	0	2	1	1	0
<i>L. melanotonus</i>	0	0	1	0	0	0	0	0	1	0	1	0	0	0	1	0	0	0	0
<i>L. chaquensis</i>	0	0	1	0	0	0	0	1	0	0	2	0	0	0	3	0&3	1&3	0&1	0
<i>L. insularum</i>	0	0	1	0	0	0	0	0	0	0	1	1	0	0	0	2	0	0	0
<i>L. pentadactylus</i>	0	0	1	0	0	2	0	1	1	1	1	1	0	0	0	2	2	0	0
<i>L. diedrus</i>	0	0	1	0	0	0	0	0	1	0	0	1	0	1	0	2	1&2	1	1
<i>L. riveroi</i>	0	0	1	0	0	0	0	1	1	0	0	1	0	1	1	3	3	0	0
<i>L. silvanimimus</i>	0	0	1	0	1	1	0	0	0	0	1	0	0	0	1	1	1	1	0
<i>A. hylaedactyla</i>	2	1	1	0	0	0	0	1	0	0	1	0	0	1	0	1	1	0	1
<i>Lith. lineatus</i>	1	0	1	1	0	0	0	0	0	0	1	0	0	1	0	0	1	1	0
<i>V. discodactylus</i>	0	0	1	0	0	0	0	1	0	0	1	0	0	1	0	2	1	1	3
<i>P. gracilis</i>	2	1	0	0	0	0	0	0	0	0	0	0	0	1	0	0	4	1	1

(See text and Heyer, 1998, for character state descriptions).

## APPENDIX 2

Molecular data matrix used for phylogenetic analysis. Regions in brackets correspond to ambiguous alignment and were not included in the analyses.

BEGINS 12S DATA	
Diedrus	[AGCGCTGAAAGATGCTGAGATGGACCCCTAAAAAGTCCTTTAACACA ] CAAAGGTTGGTCCCTAACCCCTTAAGATCAC
Riveroi	[ -GCCGTGAAAGATGCTGAGATGGACCCCTAACAAAGTCCTTTAACACA ] CAAAGGCTTGGTCCCCGGCCTTAAGATCAC
Silvani	[ -GCCGTGAAAGATGCTGAGATGGACCCCTAACAAAGTCCTTTAACACA ] CAAAGGTTGGTCTGACCTTAGAATCAC
Bufoniu	[ --CGCTGAAAGATGCTGAGATGGACCCCTAACAAAGTCCTTTAACACA ] CAAAGGTTGGTCTGACCTTAGAATCAC
Fuscuss	[ --CGCTGAAAGATGCTGAGATGGACCCCTAACAAAGTCCTTTAACACA ] CAAAGGTTGGTCTGACCTTAGAATCAC
Chaqueen	[ -GCCGTGAAAGATGCTGAGATGGACCCCTAACAAAGTCCTTTAACACA ] CAAAGGTTGGTCTGACCTTAGAATCAC
Insular	[ AGCGCTGAAAGATGCTGAGATGGACCCCTAACAAAGTCCTTTAACACA ] CAAAGGTTGGTCTGACCTTAGAATCAC
Tyloide	[ ---CTGAAAGATGCTGAGATGGACCCCTAACAAAGTCCTTTAACATA ] CAAAGGTTGGTCTGACCTTAGAATCAC
Melanon	[ --CGCTGAAAAATGCTGAGATGGACCCCTAACAAAGTCCTTTAACACA ] CAAAGGTTGGTCTGACCTTAGAATCAC
Pentada	[ -GCCGTGAAAGATGCTGAGATGGACCCCTAACAAAGTCCTTTAACACA ] CAAAGGTTGGTCTGACCTTAGAATCAC
Vanzoli	[ AGCGCTGAAAGATGCTGAGATGGACCCCTAACAAAGTCCTTTAACACA ] TAAAGGTTGGTCTGACCTTAGAATCAC
Adenhya	[ ---GCTGAAGATGCTGAGATGGACCCCTAACAAAGTCCTTTAACACA ] CAAAGGTTGGTCTGACCTTAGAATCAC
Lithody	[ ---GCTGAAGATGCTGAGATGGACCCCTAACAAAGTCCTTTAACACA ] CAAAGGTTGGTCTGACCTTAGAATCAC
Physala	[ ---GCTGAAGATGCTGAGATGGACCCCTAACAAAGTCCTTTAACACA ] CAAAGGTTGGTCTGACCTTAGAATCAC
Diedrus	TCTTACTTAACATGCAAGTCTAGCACCCTGTGTGAAAAGGCCCTTAACCTCCT-ACA-AGGGCAAGGAG
Riveroi	TCTTACTTAACCTACATGCAAGTCTAGCACCCTGTGTGAAAAGGCCCTTAACCTCCA-CTA-AGGAACAAGGAG
Silvani	TCTTACTTAACATGCAAGKCTAGCACCCTGTGTGAAAAGGCCCTTAACCTCCC-CC-TGGAGTAAGGAG
Bufoniu	TTTTTACTTAATTACATGCAAGTCTAGCACCCTGTGTGAAAAGGCCCTTAACCTCCCTAGCGGGACAAGGAG
Fuscuss	TTTTTACTTAATTACATGCAAGTCTAGCACCCTGTGTGAAAAGGCCCTTAACCCCT-AAA-AGGGACGGAG
Chaqueen	TTTTTACTTAACATGCAAGTCTAGCACCCTGTGTGAAAAGGCCCTTAACCTCC-ATT-AGGAACAAGGAG
Insular	TTTTTACTTAATTACATGCAAGTCTAGCACCCTGTGTGAAAAGGCCCTTAACCTCCCTAAGGCCCTTAAGGCCCTA-AGGAGCAAGGAG
Tyloide	TTTTTACTTAATTACATGCAAGTCTAGCACCCTGTGTGAAAAGGCCCTTAACCTCC-GTT-AGGAACAAGGAG
Melanon	TTTTTACTTAATTACATGCAAGTCTAGCACCCTGTGTGAAAAGGCCCTTAACCTCC-TTA-CGGAACAAAGGAG
Pentada	TGTTACTTAACATGCAAGTCTAGCACCCTGTGTGAAAAGGCCCTTAACCCCT-TTA-AGGGAAAGGAG
Vanzoli	TCTTACTTAACATGCAAGTCTAGCACCCTGTGTGAAAAGGCCCTTAACCTCCTAGACCCCT-CAA-AGGGAAAGGAG
Adenhya	TTTTTACTTAATTACATGCAAGTATCGGCACCCCTGTGTGAAAAGGCCCTTAATTCCT-TAT-AGGGATAAGGAG
Lithody	TTTTTCTTAACATGCAAGTATCGGCACCCCTGTGTGAAAAGGCCCTTAATCCCC-GA-TAGGATAAGGAG
Physala	TATTACTTAACATGCAAGTCTAGCACCCTGTGTGAAAAGGCCCTTAATCCCC-TCT-CGGGATAAGGAG

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Diedrus	CCGGTATCAGGGCACACCAA--AAGCCCCAACCTAGCTATGCCACACCCCACAAGGGAAACTCAGCAGTGATTAAAC
Riveroi	CCGGTATCAGGGACAAGTTTAGCCCCAACCTAGCTATGCCACACCCCACAAGGGAAACTCAGCAGTGATTAAAC
Silvani	CTGGTATCAGGGCAAAACCT-TAGCCCCAACCTAGCTATGCCACACCCCACAAGGGAAACTCAGCAGTGATTAAAC
Bufoniu	CTGGTATCAGGGCACAAACAT-TAGCCCCAACCTAGCTATGCCACACCCCACAAGGGAAACTCAGCAGTGATTAAAC
Fuscuss	CTGGTATCAGGGCACAAACAT-TAGCCCCAACCTAGCTATGCCACACCCCACAAGGGAAACTCAGCAGTGATTAAAC
Chaqueen	CTGGTATCAGGGCACAAACCTT-TAGCCCCAACCTAGCTATGCCACACCCCACAAGGGAAACTCAGCAGTGATTAAAC
Insular	CTGGTATCAGGGCACAAACATCT-TAGCCCCAACCTAGCTATGCCACACCCCACAAGGGAAACTCAGCAGTGATTAAAC
Tyloide	CTGGTATCAGGGCACAAACCTT-TAGCCCCAACCTAGCTATGCCACACCCCACAAGGGAAACTCAGCAGTGATTAAAC
Melanon	CTGGTATCAGGGCACAAATA-TAGCCCCAACCTAGCTATGCCACACCCCACAAGGGAAACTCAGCAGTGATTAAAC
Pentada	TTGGTATCAGGGCTCAAACAT-TAGCCCCACATCTCT-TAGCCCCAACCTAGCTATGCCACACCCCACAAGGGAAACTCAGCAGTGATTAAAC
Vanzoli	CCGGTATCAGGGCACATCAAATAAGCCCCAACCTAGCTATGCCACACCCCACAAGGGAAACTCAGCAGTGATTAAAC
Adenhya	CCGGTATCAGGGCACATCAAATAAGCCCCAACCTAGCTATGCCACACCCCACAAGGGAAACTCAGCAGTGATCAC
Lithody	CTGGTATCAGGGCACAAATT-TAGCCCCAACCTAGCTAGCCACACCCCACAAGGGAAACTCAGCAGAAACTCAGCAGTGATTAAAC
Physala	CTGGTATCAGGGCCCCAAAATTCT-GCCCCAAACACTAGCTATGCCACATCCACAAAGGAACACTAGCAGTGATTAAAC
Diedrus	ATTAAACATGAGCGAGCTTGATTCAAGTAAGAAAAAGAGGCCAAATCTGGTGCCTAGCCAGCCGCCGGGTAA
Riveroi	ATTGGCATGAGCGCCAGCTCGACTCAATTAAAGTAAAAGGCCAAATCTGGTGCCTAGCCAGCCGCCGGGTAA
Silvani	ATTGAATAAGGAGCAAGCTTGACTCAATTAAAGTAAAAGGCCAAATCTGGTGCCTAGCCAGCCGCCGGGTAA
Bufoniu	ATTGAATAAGGAGCAAGCTTGACTCAATTAAAGTAAAAGGCCAAATCTGGTGCCTAGCCAGCCGCCGGGTAA
Fuscuss	ATTGAATAAGGAGCAAGCTTGACTCAATTAAAGTAAAAGGCCAAATCTGGTGCCTAGCCAGCCGCCGGGTAA
Chaqueen	ATTGAATAAGGAGCAAGCTTGACTCAATTAAAGTAAAAGGCCAAATCTGGTGCCTAGCCAGCCGCCGGGTAA
Insular	ATTGAATAAGGAGCAAGCTTGACTCAATTAAAGTAAAAGGCCAAATCTGGTGCCTAGCCAGCCGCCGGGTAA
Tyloide	ATTGAATAAGGAGCAAGCTTGACTCAATTAAAGTAAAAGGCCAAATCTGGTGCCTAGCCAGCCGCCGGGTAA
Melanon	ATTGGACATAAGGAGCAAGCTTGACTCAATTAAAGTAAAAGGCCAAATCTGGTGCCTAGCCAGCCGCCGGGTAA
Pentada	ATTGAATAAGGAGCAAGCTTGACTCAATTAAAGTAAAAGGCCAAATCTGGTGCCTAGCCAGCCGCCGGGTAA
Vanzoli	ATTAAACATAAGGAGCAAGCTTGACTCAATTAAAGTAAAAGGCCAAATCTGGTGCCTAGCCAGCCGCCGGGTAA
Adenhya	ATTAAATATCAGCGAGCTTGACTCAATTAAAGTAAAATAGAGGCCGGCTAATCTGGTGCCTAGCCAGCCGCCGGGTAA
Lithody	ATTGAACATCAGCGAGCTGGATTCAAGTAAAAGTTACAGAGGCCGGCTAATCTGGTGCCTAGCCAGCCGCCGGGTAA
Physala	ATTGAACATAAGCGAGCTTGATTCAAGTAAAAGTAAATCTGGTGCCTAGCCAGCCGCCGGGTAA

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Diedrus	GCTAGGAAACAAACTGGGATTAGATAACCCCACTATGCCCTAGCCGTAAACTTAACTTACACCT-CAATCGCCCCGG	Diedrus	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTACCCCAAATCCACCTAGAGGGCCTGTCTACAAATCG
Riveroi	GCGGGGAACAAACTGGGATTAGATAACCCCACTATGCCCTAGCCGTAAACTTAACTTAC-CTCCAAATGCCAAGGG	Riveroi	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTACCCCAAATCCACCTAGAGGGCCTGTCTATAATCG
Silvani	GCTTGGGAACAAACTGGGATTAGATAACCCCACTATGCCCTAGCCGTAAACTTAACTTACAACT-TGATCGCCTGGG	Silvani	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTACCCCAAATCCACCTAGAGGGCCTGTCTATAATCG
Bufoniu	GCTAGGAACAAACTGGGATTAGATAACCCCACTATGCCCTAGCCGTAAACTTAACTTACAACT-TGATCGCCTGGG	Bufoniu	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTACCCCAAATCCACCTAGAGGGCCTGTCTATAATCG
Fuscuss	GTTAGGAACAAACTGGGATTAGATAACCCCACTATGCCCTAACCGTAAACTTAACTTACACCTTTATGCCCGGG	Fuscuss	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTACCCCAAATCCACCTAGAGGGCCTGTCTATAATCG
Chaquen	GCTTGGGAACAAACTGGGATTAGATAACCCCACTATGCCCTAGCCGTAAACTTAACTTACACCT-CAATGCCAAGGG	Chaquen	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTACCCCAAATCCACCTAGAGGGCCTGTCTATAATCG
Insular	GCTTGGGAACAAACTGGGATTAGATAACCCCACTATGCCCTAGCCGTAAACTTAACTTACACCT-CCATGCCAAGGG	Insular	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTACCCCAAATCCACCTAGAGGGCCTGTCTATAATCG
Tyloide	GCTTGGGAACAAACTGGGATTAGATAACCCCACTATGCCCTAGCCGTAAACNTTAANTTACACCT-NAATGCCNNGG	Tyloide	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTACCCCAAATCCACCTAGAGGGCCTGTCTATAATCG
Melanon	GCTTGGGAACAAACTGGGATTAGATAACCCCACTATGCCCTAGCCGTAAACTTAACTTACACCTTTATGCCCTAGGCCCTAGCCCAAGGG	Melanon	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTACCCCAAATCCACCTAGAGGGCCTGTCTATAATCG
Pentada	GCTTAGGAAACAAACTGGGATTAGATAACCCCACTATGCCCTAGCCGTAAACTTTTACACCTTACACCT-ACATGCCAAGGG	Pentada	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTACCCCAAATCCACCTAGAGGGCCTGTCTATAATCG
Vanzoli	GCTTAGGAAACAAACTGGGATTAGATAACCCCACTATGCCCTAACCGTAAACTTAACTTACACCTTACACCT-CGATGCCAAGGG	Vanzoli	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTACCCCAAATCCACCTAGAGGGCCTGTCTATAATCG
Adenhya	GCTAAGAAACAAACTGGGATTAGATAACCCCACTATGCCCTAGCCGTAAACAAATTACACCCCCAATTCGCCCCGGG	Adenhya	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTAAACACTTACAAATATCTATGCCCAAGGG
Lithody	GTCAGAGACACAAACTGGGATTAGATAACCCCACTATGCCCTAGCCGTAAACCTTACACCT-TAATGCCCAAGGG	Lithody	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTACCCCAAATCCACCTAGAGGGCCTGTCTATAATCG
Physala	GTTAAGAGATAACACTGGGATTAGATAACCCCACTATGCCCTAGCCGTAAACCTTACACCT-TAATGCCCAAGGG	Physala	AACTACGGGCAAAGCTTAAACCCAAAGGACTTGAACGGTACCCCAAATCCACCTAGAGGGCCTGTCTATAATCG

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Diedrus	ACTCTCTACT-CTAGAAGAAA-CAAAGACTA--TATGAAAC-TTAGTCTGAAGGAGGATTAGTAGTAAAAAGAA	ATCAGAATGTTCTCTTAACCCGGCACTGGGCATGTACACACNGCCCCG
Riveroi	ACTCTCTAA-A TAGAAGAAA-CGAAAGACT-TATGAAAC-CTAGTCGAAGGAGGATTAGTAGTAAAAAGGG	ATCAGAGGTCTTTTAACCCGGCACTGGGCTGTGTACACACCGCCCCG
Silvani	ACTTTCTAGT-ATAGAAGAAA-CGAAAGACTATTATGAAAC-CTGGTCGAAGGAGGATTAGTAGTAAAAAGAA	ACCAAGAGTCTTTTAACCCGGCACTGGGCTGTGTACACACCGCCCCG
Bufoniu	ACTTTCTACC-GTAGAAAAAA-CGAAAACATTATGAAAT-CTAGTCGAAGGAGGATTAGTAGTAAAAAGAA	ATCAGAGGTCTTTTAACCCGGCACTGGGCTGTGTACACACCGCCCCG
Fuscuss	ACTTTCTACC-CTAGAAAAAAACGAAAGACTACC'TATGAAAT-CTAGTCGAAGGAGGATTAGTAGTAAAAAGAA	ACCAAGAGTCTTTTAACCCGGCACTGGGCTGTGTACACACCGCCCCG
Chaqueen	ACTTTCTATT-TTAGAAGAAA-CGAAAGACTATAATGAAAT-CTAGTCGAAGGAGGATTAGTAGTAAAAAGAA	ATCAGAGGTCTTTTAACCCGGCACTGGGCTGTGTACACACCGCCCCG
Insular	ACTTTCTACG-ATAGAAAAAA-CGAAAGACTATAATGAAAC-CTAGTTAGAAGGAGGATTAGTAGTAAAAAGAA	ACTAGAGTGTCTTTTAACCCGGCACTGGGCTGTGTACACACCGCCCCG
Tyloide	ACTTTCTAAT-TTAGAAAAAA-CGAAAAGCTATTATGAAAT-CTAGCCAGAAGGAGAAATTAGTAGTAAAAAGAA	AACAGAGTGTCTTTTAACCCGGCACTGGGCTGTGTACACACCGCCCCG
Melanon	ACTTTCTAAT-TTAGAAAAAA-CGAAAAGGTATAATGAAAC-CTAACCGAAGGAGGATTAGTAGTAAAAAGAA	AACATAGTGTCTTTTAACCCGGCACTGGGCTGTGTACACACCGCCCCG
Pentada	ACTCTCTAACC-TTAGAAAAAA-CGAAAGACCACTATGAAAC-CTGGTCGAAGGAGGATTAGTAGTAAAAAGAG	AACAGAGTGTCTTTTAACCCGGCACTGGGCTGTGTACACACCGCCCCG
Vanzoli	ACTCTCTAACC-TTAGAAAAAA-CAAAAGACTACATAATGAAAT-CTAGTTAGAAGGAGGATTAGTAGTAAAAAGAA	ACCAAAAGAGTGTCTTTTAACCCGGCACTGGGCTGTGTACACACCGCCCCG
Adenhya	ACTCCCTAAAC-CTAAGGCCACA-CGAAAACACTATGAAAC-CTAACCTGTAAAGGAGGATTAGTAGTAAAAAGAA	ACCAAAAGAGTGTCTTTTAACCCGGCACTGGGCTGTGTACACACCGCCCCG
Lithody	ACTTTCTAAC-ATAGAATATA-CGAAAAGATTACTATGAAAC-CTAACCTGTAAAGGAGGATTAGTAGTAAAAAGAA	ACCAAGAGTGTCTTTTAACCCGGCACTGGGCTGTGTACACACCGCCCCG
Physala	ACTTTCTAAT-CTAGAAAATA-CTAAAGACTACCTATGAAAT-CTAGTCTGAAGGAGGATTAGTAGTAAAAAGAA	ACCAAGAGTGTCTTTTAACCCGGCACTGGGCTGTGTACACACCGCCCCG

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BEGINS 16S DATA	
Diedrus	ATAAGAGGTCCAGGCCA-GTGAC-TCT--GTTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Riveroi	ATAAGAGGTCCGGCTGCCA-GTGAC-TCT--GTTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Silvani	ATAAGAGGTCCAGCCCTGCCA-GTGAC-TCT--GTTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Bufoniu	ATAAGAGGTCCAGCCCTGCCA-GTGAC-TCT--GTTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Fuscuss	ATGAAGGTCCAGCCCTGCCA-GTGAC-TCT--GTTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Chaquen	ATAAGAGGTCCAGCCCTGCCA-GTGAC-TCT--GTTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Insular	ATAAGAGGTCCAGCCCTGCCA-GTGAC-TCT--GTTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Tyloide	ATTAAGAGGTCTAGCCCTGCCA-GTGAC-TCT--GTTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Melanon	ATAAGAGGTCCAGCCCTGCCA-GTGAC-TCT--GTTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Pentada	ATAAGAGGTCCAGCCCTGCCA-GTGAC-TCT--GTTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Vanzoli	ATGAAGGTCCAGCCCTGCCA-GTGAC-TCT--GTTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Adenhyia	ATAAGAGGTCTAGCCCTGCCA-GTGAC-ATT--GTTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Lithody	ATAAGAGGTCCAGCCCTGCCA-GTGAC-TCT--GTTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Physala	ATAAGAGGTCCAGCCCTGCCA-GTGAC-TCA--ATTCAACGGCCGGTATCCTAACCGTGCAGGTAA
Diedrus	CACTTGTTCCTAAATAAGGACTAGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA
Riveroi	CACTTGTTCCTAAATGAGGACCGGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA
Silvani	CACTTGTTCCTAAATGAGGACTAGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA
Bufoniu	CACTTGTTCCTAAATGAGGACTAGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA
Fuscuss	CACTTGTTCCTAAATGAGGACTAGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA
Chaquen	CACTTGTTCCTAAATGAGGACTAGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA
Insular	CACTTGTTCCTAAATGAGGACTAGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA
Tyloide	CACTTGTTCCTAAATGAGGACTAGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA
Melanon	CACTTGTTCCTAAATGAGGACTAGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA
Pentada	CACTTGTTCCTAAATGAGGACTAGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA
Vanzoli	CACTTGTTCCTAAATGAGGACTAGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA
Adenhyia	CACTTGTTCCTAAATGAGGACTAGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA
Lithody	CACTTGTTCCTAAATGAGGACTAGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA
Physala	CACTTGTTCCTAAATGAGGACTAGTATGAATGGOACCACGAGGTTATACTGTCTCCATCAGTAAA

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Diedrus Riveroi	ACAGGGCAATCCACTTCAAGAGCTCCTATCGACAAGTGGGTTACGACCTCGATGTTGG-ATCAGGGCATCCTAGT
Silvani	ACAGGCCAATCCACTTCAAGGCCCTATCGACAAGTGGGTTACGACCTCGATGTTGG-ATCAGGGGTGCCCCAGT
Bufoniu	ACAGGCCAATCCACTTCAAGGCCCTATCGACAAGTGGGTTACGACCTCGATGTTGG-ATCAGGGTATCCTAGT
Fuscuss	ACAGGCCAATCCACTTCAAGGCCCTATCGACAAGTGGGTTACGACCTCGATGTTGG-ATCAGGGTATCCTAGT
Chaquen	ACAGGCCAATCCACTTCAAGGCCCTATCGACAAGTGGGTTACGACCTCGATGTTGG-ATCAGGGTATCCTAGT
Insular	ACAGGCCAATCCACTTCAAGGCCCTATCGACAAGTGGGTTACGACCTCGATGTTGG-ATCAGGGTATCCTAGT
Tyloide	ACAGGCCAATCCACTTCAAGGCCCTATCGACAAGTGGGTTACGACCTCGATGTTGG-ATCAGGGTATCCTAGT
Melanon	ACAGGCCAATCCACTTCAAGGCCCTATCGACAAGTGGGTTACGACCTCGATGTTGG-ATCAGGGNNNNCCN-AGT
Pentada	ACAGGCCAATCCACTTCAAGGCCCTATCGACAAGTGGGTTACGACCTCGATGTTGG-ATCAGGGTATCCTAGT
Vanzoli	ACAGGCCAATCCACTTCAAGGCCCTATCGACAAGTGGGTTACGACCTCGATGTTGG-ATCAGGGTCCCTAGT
Adenhya	ACAGGCCAATCCACTTCAAGGCCCTATCGACAAGTGGGCTTACGACCTCGATGTTGG-ATCAGGGTACCCAGT
Lithody	ACAGGCCAATCCACTTCAAGGCCCTATCGACAAGTGGGTTACGACCTCGATGTTGG-ATCAGGGTATCCAGT
Physala	ACAGGCCAATCCACTTCAAGGCCCTATCGACAAGTGGGTTACGACCTCGATGTTGG-ATCAGGGTATCCAGT
Diedrus Riveroi	GGTGTAGCCGCTACTAAAGGTTCTGTTTGTTCAACGATTAAAAACCCCTACGTGATCTGAGTTCAAGCCGGAGTAATCC
Silvani	GGTGCAGGCCGCTACTAAAGGTTCTGTTCAACGATTAAAAACCCCTACGTGATCTGAGTTCAAGCCGGAGTAATCC
Bufoniu	GGTGCAGGCCGCTACTAAAGGTTCTGTTCAACGATTAAAAACCCCTACGTGATCTGAGTTCAAGCCGGAGTAATCC
Fuscuss	GGTGCAGGCCGCTACTAAAGGTTCTGTTCAACGATTAAAAACCCCTACGTGATCTGAGTTCAAGCCGGAGTAATCC
Chaquen	GGTGCAGGCCGCTACTAAAGGTTCTGTTCAACGATTAAAAACCCCTACGTGATCTGAGTTCAAGCCGGAGTAATCC
Insular	GGTGCAGGCCGCTACTAAAGGTTCTGTTCAACGATTAAAAACCCCTACGTGATCTGAGTTCAAGCCGGAGCAATCC
Tyloide	GGTGNAGCCGCTACTAAAGGTTCTGTTCAACGATTAAAAACCCCTACGTGATCTGAGTTCAAGCCGGAGTAATCC
Melanon	GGTGCAGGCCGCTACTAAAGGTTCTGTTCAACGATTAAAAACCCCTACGTGATCTGAGTTCAAGCCGGAGTAATCC
Pentada	GGTGCAGGCCGCTACTAAAGGTTCTGTTCAACGATTAAAAACCCCTACGTGATCTGAGTTCAAGCCGGAGTAATCC
Vanzoli	GGTGCAGGCCGCTACTAAAGGTTCTGTTCAACGATTAAAAACCCCTACGTGATCTGAGTTCAAGCCGGAGCAATCC
Adenhya	GGTGCAGGCCGCTACTAAAGGTTCTGTTCAACGATTAAAAACCCCTACGTGATCTGAGTTCAAGCCGGAGTAATCC
Lithody	GGTGCAGGCCGCTACTAAAGGTTCTGTTCAACGATTAAAAACCCCTACGTGATCTGAGTTCAAGCCGGAGTAATCC
Physala	GGTGCAGGCCGCTACTAAAGGTTCTGTTCAACGATTAAAAACCCCTACGTGATCTGAGTTCAAGCCGGAGTAATCC

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<i>Diedrus</i>	AGGTCAAGTTCTATCTATAAAGAGTTTCTCCAGTACGAAAGGACCGAAAAAACATGGCCAATGCCCAATGGCCCCAGTAAGCC
<i>Riveroi</i>	AGGTCAAGTTCTATCTATAAAGAGCTTTCTAGTACGAAAAGGACCGAAAAAACATGGCCCATGGCTAACATGCAAGCC
<i>Silvani</i>	AGGTCAAGTTCTATCTATAAAGAGTTTCTAGTACGAAAAGGACCGAAAAAACATGGCCCATACTTCATGCAAGCC
<i>Bufoniu</i>	AGGTCAAGTTCTATCTATAAAGAGCTTTCTAGTACGAAAAGGACCGAAAAAACATGGCCCATACCTTAAGCAAGCC
<i>Fuscuss</i>	AGGTCAAGTTCTATCTATAAAGAGCTTTCTAGTACGAAAAGGACCGAAAAAACATGGCCCATAGCAAGCC
<i>Chaquen</i>	AGGTCAAGTTCTATCTATAAAGAGCTTTCTAGTACGAAAAGGACCGAAAAAACATGGCCCATACTTCATGCAAGCC
<i>Insular</i>	AGGTCAAGTTCTATCTATAAAGAGCTTTCTAGTACGAAAAGGACCGAAAAAACATGGCCCATACTATATGCAAGCC
<i>Tyloide</i>	AGGTCAAGTTCTATCTATAAAGAGTTTCTAGTACGAAAAGGACCGAAAAAACATGGCCCATAGCAAGCC
<i>Melanon</i>	AGGTCAAGTTCTATCNATAAAGAGTTTCTAGTACGAAAAGGACCGAAAAAACATGGCCCATAGCAAGCC
<i>Pentada</i>	AGGTCAAGTTCTATCTATAAAAGCTTTCTAGTACGAAAAGGACCGAAAAAACATGGCCCATACTTCATGCAAGCC
<i>Vanzoli</i>	AGGTCAAGTTCTATCTATAAAGAGCTTTCTAGTACGAAAAGGACCGAAAAAACATGGCCCATATGCAAGCC
<i>Adenhy</i>	AGGTCAAGTTCTATCTATAAAGAGCTTTCTAGTACGAAAAGGACCGAAAAAACATGGCCCATAGCTTC-AATAAGCC
<i>Lithody</i>	AGGTCAAGTTCTATCTATAAAGAGCTTTCTAGTACGAAAAGGACCGAAAAAACATGGCCCATTTTATTACAACC
<i>Physala</i>	AGGTCAAGTTCTATCTATAAAGAGCTTTCTAGTACGAAAAGGACCGAAAAAACATGGCCCATATTTTATATAAGCC
<i>Diedrus</i>	ATACAAACCTATTATG-ACACAT
<i>Ri Veroi</i>	ATACAAATTATG-ATACAC
<i>Silvani</i>	ATACAACTAAATTATGACACAC
<i>Bufoniu</i>	GTAAGCAACCAATTATG-ACACAGC
<i>Fuscuss</i>	GTAACAACCAACTTATG-ACATAG
<i>Chaquen</i>	ATAACAGATAATTATG-ACACAAAC
<i>Insular</i>	ATAACAACTAATTATG-ACACAAAC
<i>Tyloide</i>	ATAGCAACCTATTATG-ACTTAAC
<i>Melanon</i>	ATAACGCTCAATTATG-ACTAAAC
<i>Pentada</i>	ATAACAGCCATTATG-ACATAAC
<i>Vanzoli</i>	ATAATAACCTTATTATG-ACCAAAT
<i>Adenhy</i>	ATACCAATT-ATTATG-AATTAT
<i>Lithody</i>	ATTCTAATTAACTTTG-ACTTAAC
<i>Physala</i>	ATAGT--CTAATTATG-TTTATAC

