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# The Status of *Leptodactylus Validus* as a Single Taxon and Its Relationship to *Leptodactylus Pallidirostris*

Kenneth Christopher Yanek

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THE STATUS OF *LEPTODACTYLUS VALIDUS* AS A SINGLE TAXON AND  
ITS RELATIONSHIP TO *LEPTODACTYLUS PALLIDIROSTRIS*

Kenneth Christopher Yanek

Master of Science in Biology

University of Richmond, 2004

Thesis Director: Dr. Rafael O. de Sá

**Abstract.** *Leptodactylus validus* exhibits an unusual distribution inhabiting Trinidad, Tobago, and the Lesser Antilles, but not the mainland of South America. This distribution is inconsistent with other distribution patterns observed for these islands. Although slight variation in adult morphology has been observed among the island populations, call data suggest the presence of a single species. Calls of *Leptodactylus pallidirostris* from Venezuela suggested that this taxon might be conspecific with *L. validus*. Herein sequence data from the 12S and 16S mt rRNA genes were collected and analyzed in order to address: 1.) whether more than one species is represented within *L. validus*, and 2.) to determine if *L. pallidirostris* is conspecific with *L. validus*. Independent and combined phylogenetic analyses were conducted using maximum parsimony,

maximum likelihood, and Bayesian analysis. Results obtained indicate that *L. validus* represents a single species throughout its distribution and is conspecific with *L. pallidirostris*.

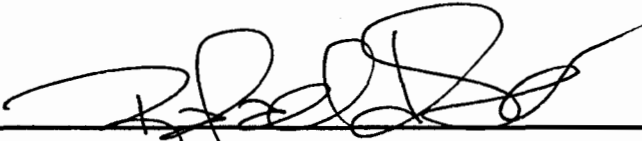
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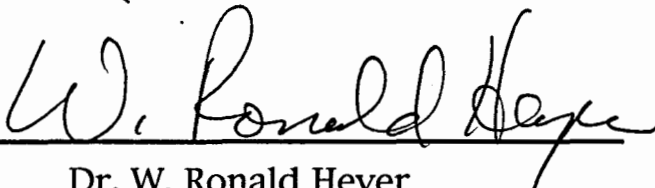
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I certify that I have read this thesis and find that, in scope and  
quality, it satisfies the requirements for the degree of Master of  
Science.




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Dr. Rafael O. de Sá, Thesis Advisor



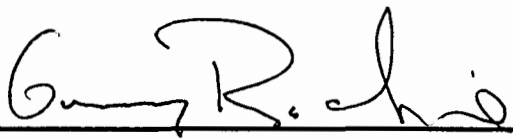
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Dr. W. Ronald Heyer



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Dr. W. John Hayden



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THE STATUS OF *LEPTODACTYLUS VALIDUS* AS A SINGLE TAXON AND  
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By

KENNETH CHRISTOPHER YANEK

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*To my parents Kathy and Kenneth T. Yanek*

## Introduction

The frog genus *Leptodactylus* (Anura: Leptodactylidae: Leptodactylinae) consists of approximately 70 species distributed throughout Latin America including portions of the West Indies extending through the lowlands of Mexico and southern Texas (Heyer, 1969a; Frost, 2002). Four species groups are currently recognized within this genus: '*fuscus*', '*melanonotus*', '*ocellatus*', and '*pentadactylus*' (Heyer, 1969b, 1974; Maxon and Heyer, 1988; Frost, 2002). Heyer (1969b) originally described the evolutionary relationships among members of the four species groups as representing a transition from aquatic to terrestrial modes of life. This is exemplified by the adaptive morphology, behavior, and the modes of reproduction employed by members within the different groups.

*Leptodactylus validus* is a cryptically colored frog belonging to the *melanonotus* species group. Members of this group possess extensive toe fringes as adults and dorsolateral folds, if present, are weakly defined with a warty texture (Heyer, 1970). Males possess cornified thumb spines that are used primarily in combat for

defending territories (Shine, 1979) but do not have chest spines (Heyer, 1969; 1970). Species in the *melanonotus* group construct foam nests that are placed on the surface of water. In a review of the relationships of the *melanonotus* species group, Heyer (1970) used a set of 34 morphological characters to attempt to explain their interspecific relationships and distribution patterns. That analysis resulted in the placement of *Leptodactylus validus* in the synonymy of *L. wagneri*. Subsequently, specimens from additional localities became available and *L. validus* was again recognized as a valid taxon, and included in the *podicipinus-wagneri* complex (Heyer, 1994). This complex currently consists of 13 species whose interspecific relationships are not well understood.

*Leptodactylus validus* was first described by Garman (1887) from Kingston, St. Vincent based on a series of specimens with no designation of a single specimen as the holotype. Subsequently, Heyer (1970) designated MCZ 2185 from Kingston St. Vincent as the lectotype of *Leptodactylus validus* Garman. The distribution of *L. validus* includes Trinidad, Tobago, and the Lesser Antillean islands of Bequia, Grenada, and St. Vincent. Males range in size between

30-45 mm SVL while females range between 38-56 mm (Heyer, 1994; Murphy, 1997). *Leptodactylus validus* is distinguished from other sympatric members of the genus by its lack of long dorsolateral folds and, in some cases, the presence of a discernable lip stripe extending posteriorly from the eye to the angle of the mouth (Heyer, 1994). A terrestrial species, *L. validus* is often found among leaf litter or debris in or near forests, along heavily shaded streams and gullies, within caves, or roadside ditches where adequate cover is available (Johnson, 1946; Kenny, 1969; Downie, 1996; Murphy, 1997). Breeding occurs during the heavy summer rains and mating takes place in ponds, drainage ditches, or temporary flooded forest areas (Kenny, 1969; Downie, 1996; Murphy, 1997; Germano et al., 2003). Eggs are deposited in foam nests produced from oviduct secretions by the female. Foam nests are constructed along the edges of pools but also may be free floating and subsequently covered by dead vegetation such as leaves or sticks (Kenny, 1969; Downie, 1996). Within 2.5-5 days, the larvae hatch and remain in contact with the mucus at the base of the nest (Kenny, 1969; Downie, 1996). Soon after, the tadpoles

form tight shoals and spend much of their time near the surface (Downie, 1996). Parental care in *L. validus* has been observed with females remaining with the nests and shoals of tadpoles for some time after hatching (Downie, 1996). Metamorphosis usually occurs within 8 weeks (Kenny, 1969). The advertisement call of *L. validus* has a broadcast frequency range of 1300-3500 Hz with maximum energy between 2300-3500 Hz. It usually consists of two notes lasting 0.03-0.06 seconds in duration at a rate of 1.1-1.9 calls per second (Heyer, 1994).

Slight morphological variation exists between adult populations from the Lesser Antilles and those from Trinidad and Tobago. Additionally, the descriptions of *L. validus* tadpoles from Trinidad, Tobago, and St. Vincent differ drastically (Heyer, 1994), however the number of larvae in collections is limited. Heyer (1994) suggested that the tadpole from Trinidad described by Kenny (1969), may have been misidentified, and might belong to a genus other than *Leptodactylus*. Furthermore, the identification of larval specimens from St. Vincent and Tobago were based upon known adult occurrences in the areas where they were collected, not

on positive identification of an adult female associated with them, or from a developmental series. Therefore, the validity of these descriptions is also questionable (Heyer, 1994). If the larval specimens from the three islands indeed correspond to *L. validus*, enough variation exists to suggest that *L. validus* may represent more than one taxon, with *L. validus* occurring on the Lesser Antilles and a second, distinct, and closely related species endemic to Trinidad and Tobago (Heyer, 1994).

Available call data for *L. validus* from Trinidad and Tobago is similar to calls analyzed from Venezuelan populations of *L. pallidirostris*, another species within the *L. podicipinus-wagneri* complex. This suggests that at least some populations of *L. pallidirostris* may be conspecific with *L. validus*, extending the distribution of *Leptodactylus validus* to include the mainland of South America. *Leptodactylus pallidirostris* is distributed throughout Venezuela, Guyana, Suriname, French Guiana, and northern Brazil, however the only available call data comes from Brazil and Venezuela (Heyer, 1994).



Heyer (1994) found extensive interspecific and intraspecific morphological variation within the *Leptodactylus podicipinus-wagneri* complex; this variation limits the use of morphology to decipher relationships. The analysis of mitochondrial DNA (mtDNA) sequence data has become a powerful tool frequently implemented to depict evolutionary relationships in systematics. MtDNA is useful because it is maternally inherited, has a relatively rapid rate of evolution, and its genes are relatively easy to amplify and sequence (Awise, 1986, 1994, 2000; Wilson et al., 1985; Moritz et al., 1987). Additionally, differing natural selection pressures on a population's external phenotype may have a significant impact on a phylogeny obtained using morphological characters, while this is less likely to occur using mtDNA sequence data (Wüster et al., 2002). The mitochondrial ribosomal RNA (mt rRNA) genes are particularly useful in addressing divergences that have occurred within the past 100 million years among closely related taxa.

Herein sequence data from the small (12S) and large (16S) subunit mt rRNA genes were collected and analyzed in order to address: 1.) whether more than one species is represented within *L.*

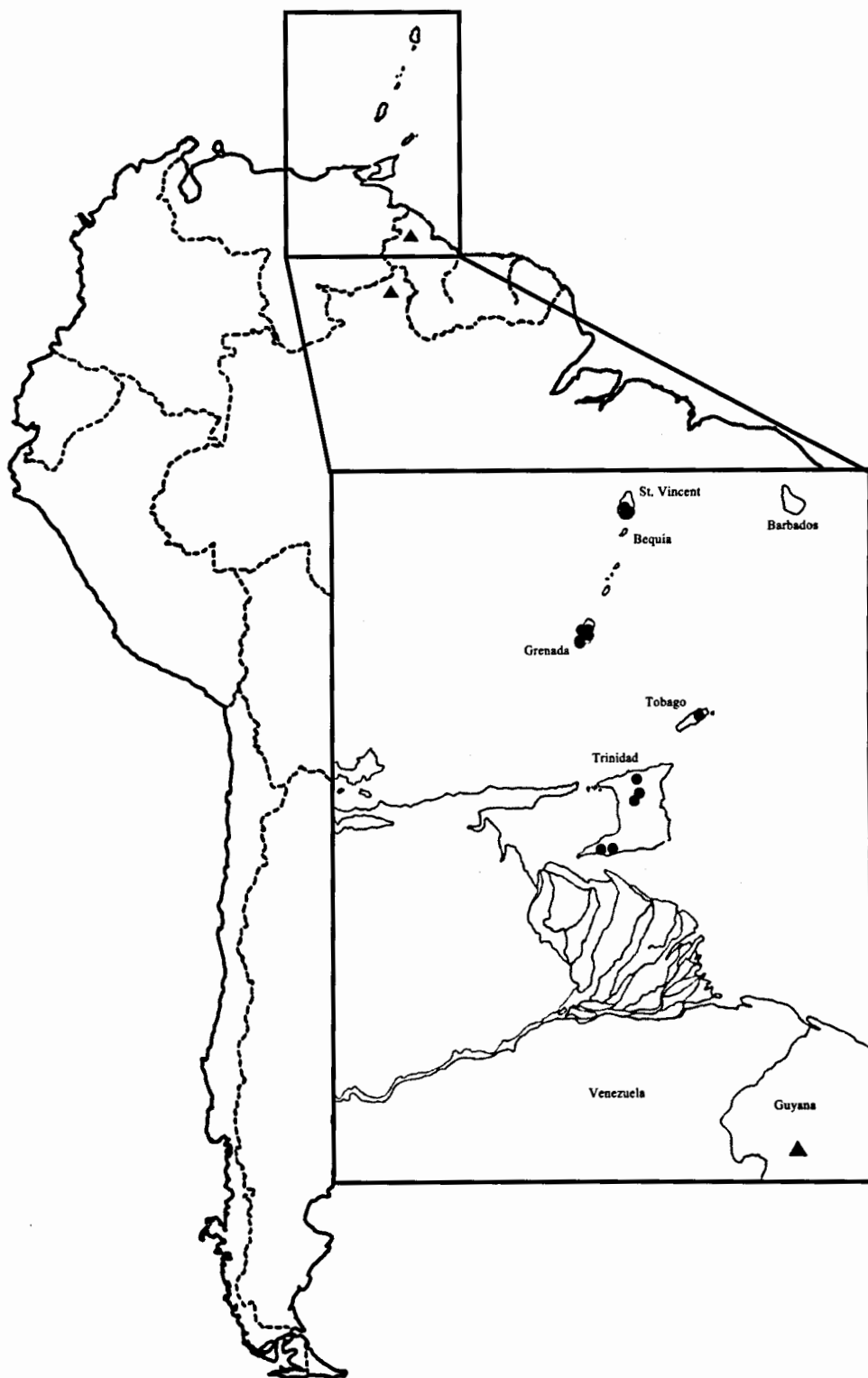
*validus*, and 2.) to determine if at least some populations currently recognized as *L. pallidirostris* are conspecific with *L. validus*.

## Materials & Methods

Frozen tissue samples (liver and muscle) of *Leptodactylus validus* from Trinidad (n=15), Tobago (n=2), and the Lesser Antillean Islands of St. Vincent (n=13), and Grenada (n=20) in addition to *L. pallidirostris* samples (from Brazil n=1 and Guyana n=1) were included in this study (Fig. 1). Samples from two additional taxa belonging to the *L. podicipinus-wagneri* complex were also included as ingroup samples: *L. podicipinus* (from Brazil n=4) and *L. wagneri* (from Brazil n=1 and from Ecuador n=2). *Leptodactylus chaquensis* (n=1) (*L. ocellatus* species group) and *L. knudseni* (n=1) (*L. pentadactylus* species group) were used as outgroups. Locality data for the samples examined are provided in Appendix I.

*DNA isolation, amplification, and purification.* — DNA was isolated following protocols of Hillis et al. (1996). Tissues were mechanically broken down and incubated at 55°C with proteinase K (10mg/ml), 10% SDS, and STE. Following incubation, DNA was extracted twice with Phenol: Chloroform: Isoamyl alcohol (25:24:1) and two subsequent extractions were performed with chloroform.

Figure 1. Map of localities where samples of *L. pallidirostris* (triangles) and *L. validus* (circles) used in this study were collected.



DNA was precipitated with cold absolute ethanol and 2M NaCl. The supernatant was discarded and the pellet resuspended in 1X TE buffer (pH 7.5). Isolated DNA samples (5 $\mu$ m) were run in 0.8% agarose gels, at 50v, followed by staining with ethidium bromide, and illumination under UV light to assess DNA quality. Fragments of mitochondrial DNA (mtDNA) from the 12S and 16S mt rRNA genes were amplified using polymerase chain reaction (PCR; Palumbi, 1996) in an MJ Research PTC-200 thermocycler and a Stratagene SCS-2 temperature cycler. Double stranded amplifications of nearly the complete 12S (~880 base pairs) and 16S (~1,500 bp) rRNA genes were amplified using four sets of primers (Goebel et al., 1999). Primer sequences are presented in Table I. PCR reactions for the first ~570 bp of the 12S gene were performed using the primers 12S tphef and 12S Rds1 with JumpStart Taq DNA Polymerase (Sigma). PCR reactions consisted of a 2.5-min denaturation at 94°C, 1-minute of annealing at 55°C, and a 2-min extension at 72°C, followed by 30 cycles of a 1-min denaturation at 94°C, 1 minute of annealing at 55°C, and an extension period of 1.5-min at 72°C. A second set of primers (12SA-L and 12SB-H), were

used in conjunction with PCR Master Mix (Promega) to amplify a 12S gene fragment of ~390 bp in length, beginning approximately 80 bp prior to the end of the region amplified by the 12S tphef/Rds1 primers. The conditions under which PCR reactions were conducted differed from those described above only by a lower annealing temperature (53°C).

The primer 12L13 (Feller and Hedges, 1998), which attaches to a region near the end of the 12S rRNA gene, and Hedges 16H10 (Hedges, 1994) were used to amplify a segment of the 16S rRNA gene approximately 1,030 bp in length using Master Mix (Promega). PCR conditions consisted of a 2-minute denaturation at 94°C, a 1-min annealing period at 50°C and a 72°C extension for 1.5-min, followed by 34 cycles of a 1-min denaturation at 94°C, 1-min of annealing at 50°C, and a 1.5-min extension at 72°C. An additional 550 bp of the 16S gene were amplified in both directions using the primers 16SaR-L and 16Sd with Master Mix (Promega), which overlapped the end of the L13/H10 fragment by approximately 70 bp. Reaction conditions were identical to those implemented for 12L13/Hedges 16H10. Amplified products were purified using the

Wizard Plus Miniprep DNA Purification System (Promega) and suspended in 1X TE buffer.

*Sequencing.* — Purified templates were sequenced using SequiTherm Excel II DNA sequencing kits (Epicentre) in an MJ Research PTC-200 thermocycler. Single-stranded sequencing reactions were performed using primers labeled with an infrared fluorescent dye (5' IRD800; Li-Cor). The primers used for sequencing reactions were identical in sequence to those used for amplification except 16Sd, which was replaced by the primer 16SbR-H (Palumbi et al., 1991, Table I). Sequencing reactions consisted of a 2.5-min denaturation at 95°C, followed by 30 cycles of a 30 second denaturation at 95°C, 30 seconds of annealing at 58°C, and 30 seconds of extension at 70°C. Reaction products for gene fragments < 650 bp in length were run on 44 cm long 6% acrylamide gels, whereas those exceeding 650 bp were run on 66 cm long 4% acrylamide gels. Gels were .25 mm in thickness and sequences were collected using a Li-Cor 4000L automated DNA sequencer.

*Sequence Alignment.* — Image data from each single strand sequence along with the chromatographs constructed by the



BaseImagIR Ver.4.2 software (Li-Cor Biotechnology Division) were imported into the software program AlignIR (Li-Cor Biotechnology Division) and aligned with their complimentary sequence. For each strand, bands from the aligned image files and their corresponding chromatographs were visually inspected for mismatches and corrected. Amended sequences from all samples were compiled into a data matrix and aligned with ClustalX (Thompson et al., 1997) using the multiple alignment option. Alignment ambiguities were improved manually considering the criterion of parsimony when possible and through inference of secondary structure (Kjer, 1995; Hickson et al., 1996).

*Phylogenetic analyses.* — Phylogenetic analyses were inferred for both the independent as well as the combined data sets using PAUP\* 4.0b10 (Swofford, 2002). In order to reduce computation time, groups of samples sharing identical sequences were represented by a single sample. Pair-wise genetic distances were calculated under the general time reversible model with gamma distributed rate variation. Saturation of nucleotide substitutions among samples in an alignment can affect the estimation of

evolutionary distances and potentially result in misleading tree topologies (Swofford et al., 1996; Page and Holmes, 1998; Nei and Kumar, 2000; de Peer et al., 2002). To investigate levels of substitutional saturation, pair-wise uncorrected  $p$ -distances were plotted against corrected pair-wise genetic distances using general time reversible distances with and without gamma correction for variable sites, (see below) and examined for departure from a linear relationship. Additionally, the high rate of transitional mutations relative to transversions that occur in mitochondrial genes can result in saturation of transitional sites (Brown et al., 1982; Moriyama and Powell, 1997; Wakely, 1996; de Peer et al., 2002). To assess levels of saturation for transitional substitutions, pair-wise uncorrected  $p$ -distances were plotted against Kimura 2-parameter (K2p) distances. The extent to which scatter plots deviate from the  $x=y$  line indicates the relative degree of saturation.

Maximum parsimony (MP) analyses were performed under heuristic algorithms with accelerated character transformation optimization (ACCTRAN) and delayed character transformation optimization (DELTRAN) using the tree-bisection reconnection (TBR)

branch swapping option. Bases were treated as unordered character states; gaps were treated alternatively as missing data or as a fifth character state, considering that insertions or deletions (indels) also represent informative evolutionary changes (Simmons and Ochoterena, 2000; Simmons et al., 2001). Weighted analyses were performed in which transversional (tv) changes were assigned weights twice that of transitions (ti). Using the 12S and 16S rRNA secondary structure models for *Xenopus laevis* (Cannone et al., 2002), stem and loop positions were defined and additional MP and maximum likelihood (ML) analyses were performed on all three datasets assigning loop positions twice the weight as stem positions. In instances where more than one most parsimonious tree was obtained, strict and 50% majority rule (50% MR) consensus trees were constructed. Statistical stability of internal branches was assessed via nonparametric bootstrapping (Felsenstein, 1985) based on 1000 pseudoreplicates (50% MR, heuristic search).

The program Modeltest (Ver. 3.06) (Posada and Crandall, 1998) was used to select the best-fitted model of sequence evolution for each dataset, as well as likelihood parameter settings to be

implemented in ML (Fisher, 1922; Felsenstein, 1981) analyses using PAUP\*. Hierarchical Likelihood Ratio Tests (hLRTs) resulted in the selection the Tamura–Nei model (1993) with among-site rate heterogeneity (TrN+G) for all datasets. Using Akaike Information Criterion (AIC) (Bozdogan, 1987), Modeltest selected the General Time Reversible Model (Rodriguez et al. 1990) with invariant sites and gamma-distributed rate heterogeneity (GTR+I+ $\Gamma$ ) for the 12S and combined datasets, while the General Time Reversible Model with gamma-distributed rate variation across sites (GTR+ $\Gamma$ ) was selected for the 16S dataset. For ML analyses heuristic searches were conducted using TBR branch swapping and nonparametric bootstrapping (100 pseudoreplicates, 50% majority rule). Bootstrap values 75% and greater were considered strongly supported, between 55% and 74% weak support, and values below 55% were considered low or poorly supported.

Additional likelihood analyses were performed based on Bayesian inference using MrBayes (Ver. 3.0b4) (Huelsenbeck and Ronquist. 2001). Bayesian inference uses Markov Chain Monte Carlo (MCMC) algorithms to estimate the posterior probabilities of

the best set of trees based on the evolutionary model specified. The model of sequence evolution employed for each dataset was selected using Modeltest under AIC. The number of substitution types was set to 6, enabling the rates to vary, thus being subject to the constraint of time-reversibility (Tavare, 1986). Seven simultaneous MCMC chains were run in order to determine the number of samples to discard based on convergence of log-likelihood values. Analyses were initiated using randomly selected starting trees, and topologies were sampled every 10 generations for  $2.0 \times 10^6$  generations. The resulting 50% majority-rule consensus tree was rooted using the outgroup taxa in PAUP\*. For the Bayesian analyses, credibility values for a clade were considered statistically significant when posterior support values were  $\geq 99\%$ .

## Results

In all analyses using the 12S, 16S, as well as the combined datasets, *Leptodactylus validus* and *L. pallidirostris* formed a monophyletic group. *Leptodactylus wagneri* samples formed a separate clade, as did the *L. podicipinus* samples. Additionally, pair-wise genetic distances were < 1% among the *L. validus* and *L. pallidirostris* samples in all datasets. In instances where separate analyses yielded identical tree topologies for a specified dataset, bootstrap support values for the depicted clades were within 11% of those indicated in the figures.

### 12S rRNA gene

*Sequence divergence.*— Analysis of the *Leptodactylus validus* sequences grouped the 50 individuals into five separate haplotypes (Table II). Among the haplotypes scored, five (0.6%) sites were variable within *L. validus*. Haplotype A includes samples from Grenada (n=9), St. Vincent (n= 8), Trinidad (n=1), as well as the Tobago samples (n=2). Haplotype B includes samples from Grenada (n=11), St. Vincent (n=5), and Trinidad (n=11). Haplotypes C, D,

and E each consist of a single sample from Trinidad. The aligned 828 bp fragment of the 12S rRNA gene (Appendix II) correspond to the homologous region between positions 2205–2904 of the *Xenopus laevis* mitochondrial genome (Roe et al., 1985; Gen-Bank no.: M10217), however, adjustments in the numbering scheme were made to incorporate the more accurate model as described by (Dunon-Bluteau and Brun, 1986). Base frequencies for the ingroup and outgroup taxa are presented in Table III. Corrected pair-wise genetic distances among *L. validus* samples were between 0–0.49%. The two *L. pallidirostris* samples differed by 0.25%. Genetic distances between the *L. pallidirostris* and *L. validus* samples ranged from 0.12–0.62%. Genetic distances among samples of *L. podicipinus* were between 0.12–0.88% and ranged between 10.51–11.15% when compared with *L. validus/pallidirostris*. Distances among samples of *L. wagneri* were 0.24–1.52% and 5.04–5.72% between *L. validus/pallidirostris*. Distances between *L. validus/pallidirostris* and the outgroups ranged between 12.19–12.81% compared with *L. chaquensis* and 16.93–17.5% with *L. knudseni*. Corrected pair-wise genetic distances are presented in

Table IV and absolute base differences among samples are provided in Table V. Plots of pair-wise uncorrected p-distances versus K2p distances were nearly linear, indicating a low degree of transitional saturation (Fig. 2a). Similarly, the comparison of uncorrected p-distances with corrected GTR divergences did not reveal saturation of base substitutions (Fig. 2b). However, in the comparison of uncorrected p-distances with corrected GTR+ $\Gamma$  distances, the scatter plot points began to depart from the diagonal at higher levels of sequence divergence (Fig. 2c).

*Maximum parsimony analyses.* — Maximum parsimony analysis of the 12S data considering gaps as missing data resulted in thirty equally parsimonious trees 220 steps long ( $L=220$ ) with consistency indices of 0.854 ( $CI=0.854$ ). Of the 828 aligned bp sequenced, 159 (19.2%) characters were variable and 102 (12.3%) were parsimony-informative. In all trees, *L. pallidirostris* and *L. validus* formed a monophyletic group with 100% support in the strict consensus tree (Fig. 3). Within this clade, a subclade consisting of haplotypes B and D was recovered with poor support. Relationships among the other samples were unresolved. An



Figure 2. Plots depicting relative rates of (a) transitional saturation and (b) substitutional saturation using pair-wise genetic distances corrected under the GTR and (c) GTR+ $\Gamma$  evolutionary models among 12S gene sequences.

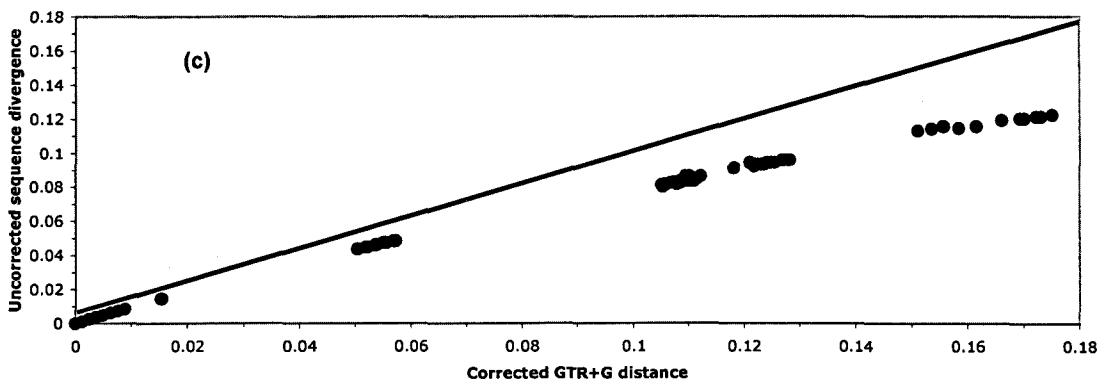
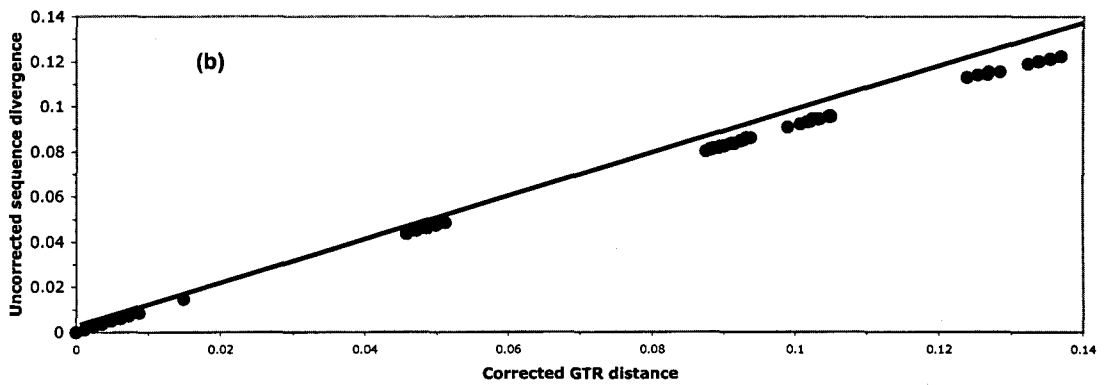
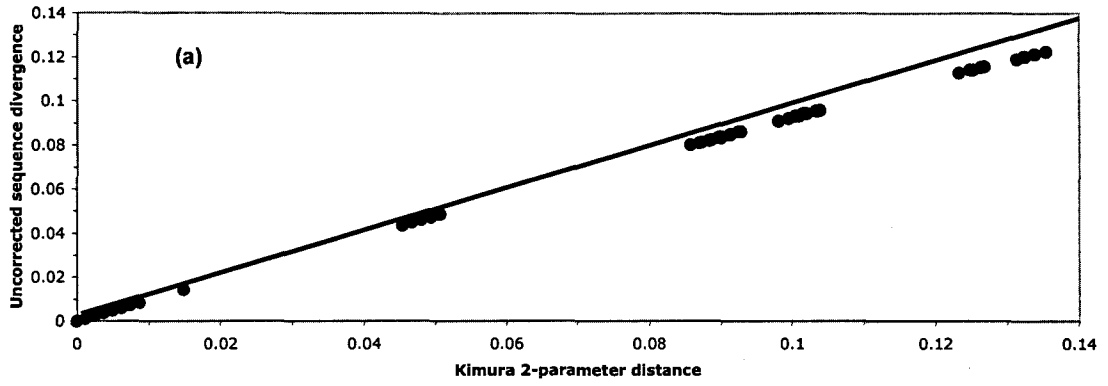
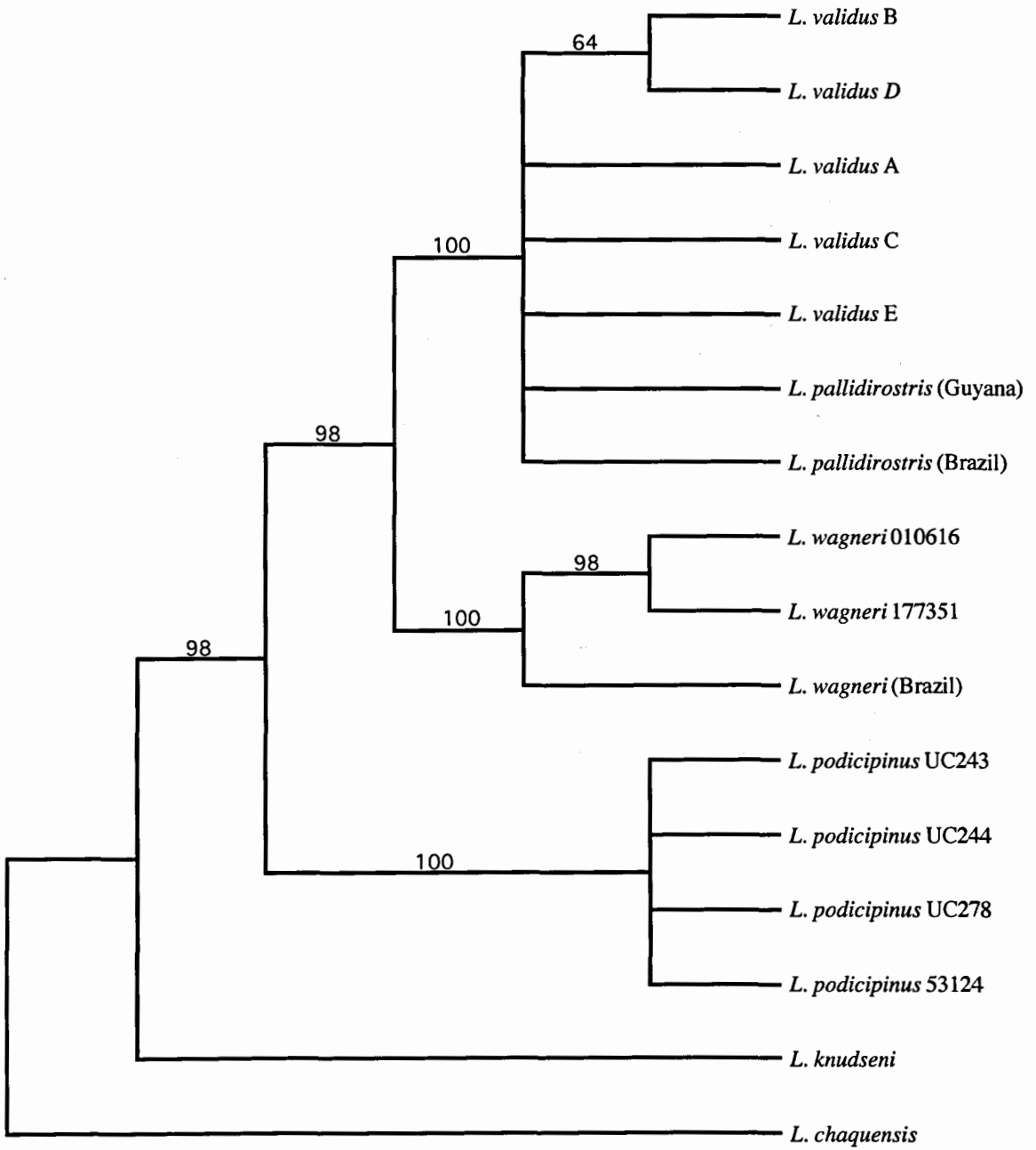


Figure 3. Consensus tree (50% MR) topology obtained in all MP analyses, unweighted ML analysis under the TrN+ $\Gamma$  evolutionary model, and both weighted ML analyses for the 12S gene. Bootstrap values shown above branches correspond to those observed in the MP analysis considering gaps as missing data. See text for clade descriptions.



analysis considering gaps as a fifth character state recovered thirty minimum-length trees (L =226; CI=0.858; 162 [19.6%] variable characters and 107 [12.9%] parsimony-informative). The strict consensus tree was identical to the tree presented in Fig. 3.

*Weighted maximum parsimony analyses.* — An analysis weighting transversions to transitions (tv:ti, 2:1), recovered thirty equally parsimonious trees (L=288; CI=0.854; 159 [19.2%] variable characters and 103 [12.4%] parsimony-informative). Weighting loop positions (n=380, weight=2) to stem positions (n=448, weight=1) and considering gaps as missing data, resulted in thirty minimum-length-trees (L=350; CI=0.843; 159 [19.2%] variable characters and 102 [12.3%] parsimony-informative). Using gaps as a fifth character state under the same weighting scheme, thirty equally parsimonious trees were obtained (L=362; CI=0.848; 162 [19.6%] variable characters and 107 [12.92] parsimony-informative). The strict consensus trees obtained in all weighed analyses shared topologies identical to that of the unweighted tree (Fig. 3).

*Maximum likelihood analyses.* — Maximum likelihood analysis of the 12S dataset implemented under the TrN+G model parameters

resulted in a bootstrap 50% Majority Rule (MR) consensus tree with a topology identical to those obtained in the MP analyses (Fig. 3). The 50% MR consensus tree obtained using the GTR+I+ $\Gamma$  model parameters resulted in a similar tree, in which a second subclade consisting of the *L. pallidirostris* specimens was present with low support (Fig. 4). ML analyses considering secondary structure and weighting loop:stem positions (2:1), using both evolutionary models resulted in bootstrap 50% MR consensus trees identical to the tree presented in Fig. 3.

*Bayesian analysis.* — A Bayesian analysis was performed using the GTR+I+ $\Gamma$  model of sequence evolution. Convergence of the Log likelihood values among the seven MCMC chains occurred within 15,000 generations of sampling, consequently the burn-in value was set to discard the first 1,500 trees sampled. The consensus tree obtained by MrBayes (Fig. 5) was similar, but slightly better resolved than those obtained in the MP and ML analyses. *Leptodactylus validus* and *L. pallidirostris* formed a monophyletic group with 100% support. Furthermore, *L. pallidirostris* and *L. validus* samples formed exclusive monophyletic groups, however these clades had

Figure 4. Unweighted 12S ML tree obtained using the GTR+I+ $\Gamma$  evolutionary model. Bootstrap values are shown above branches. See text for clade descriptions

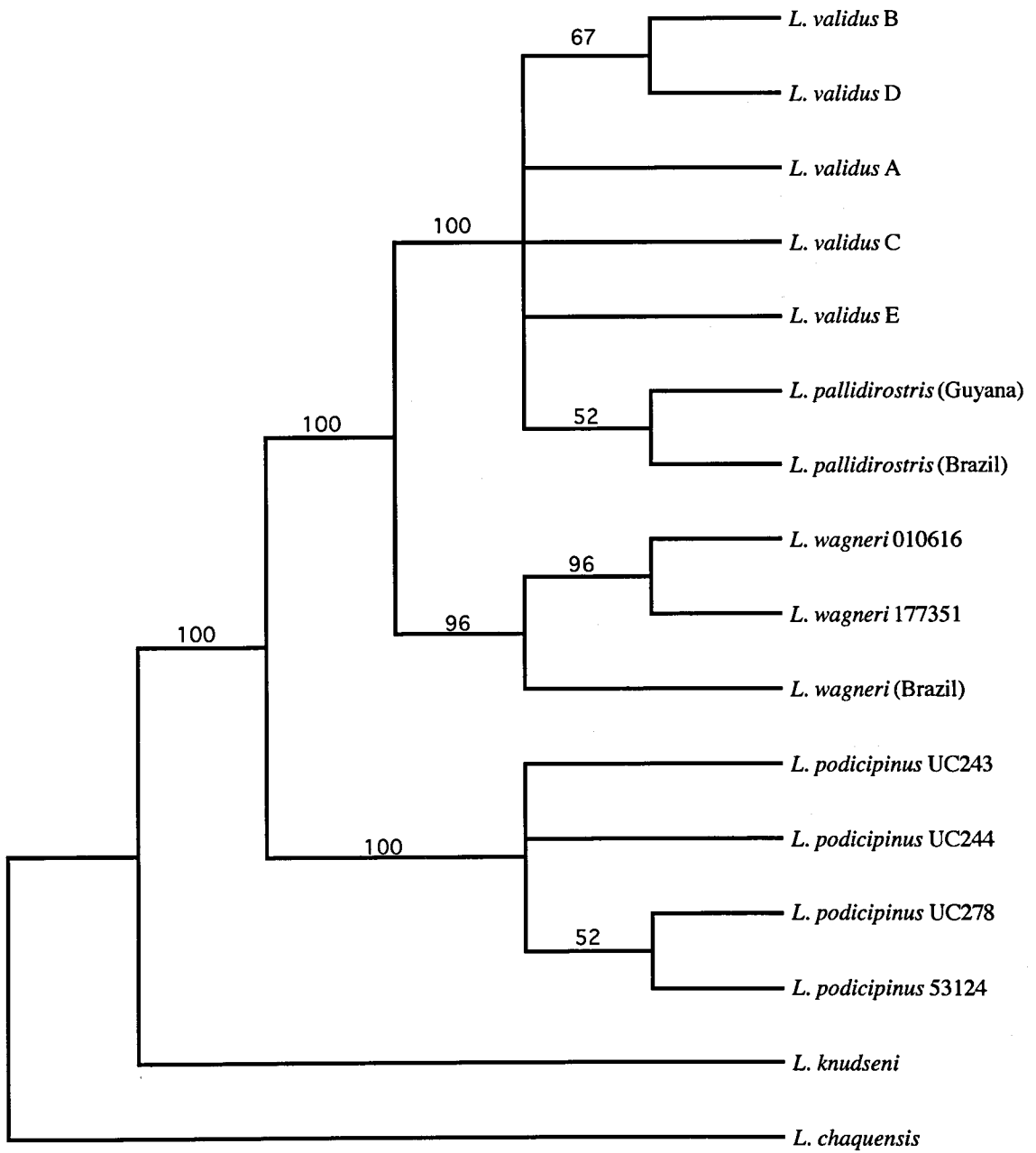
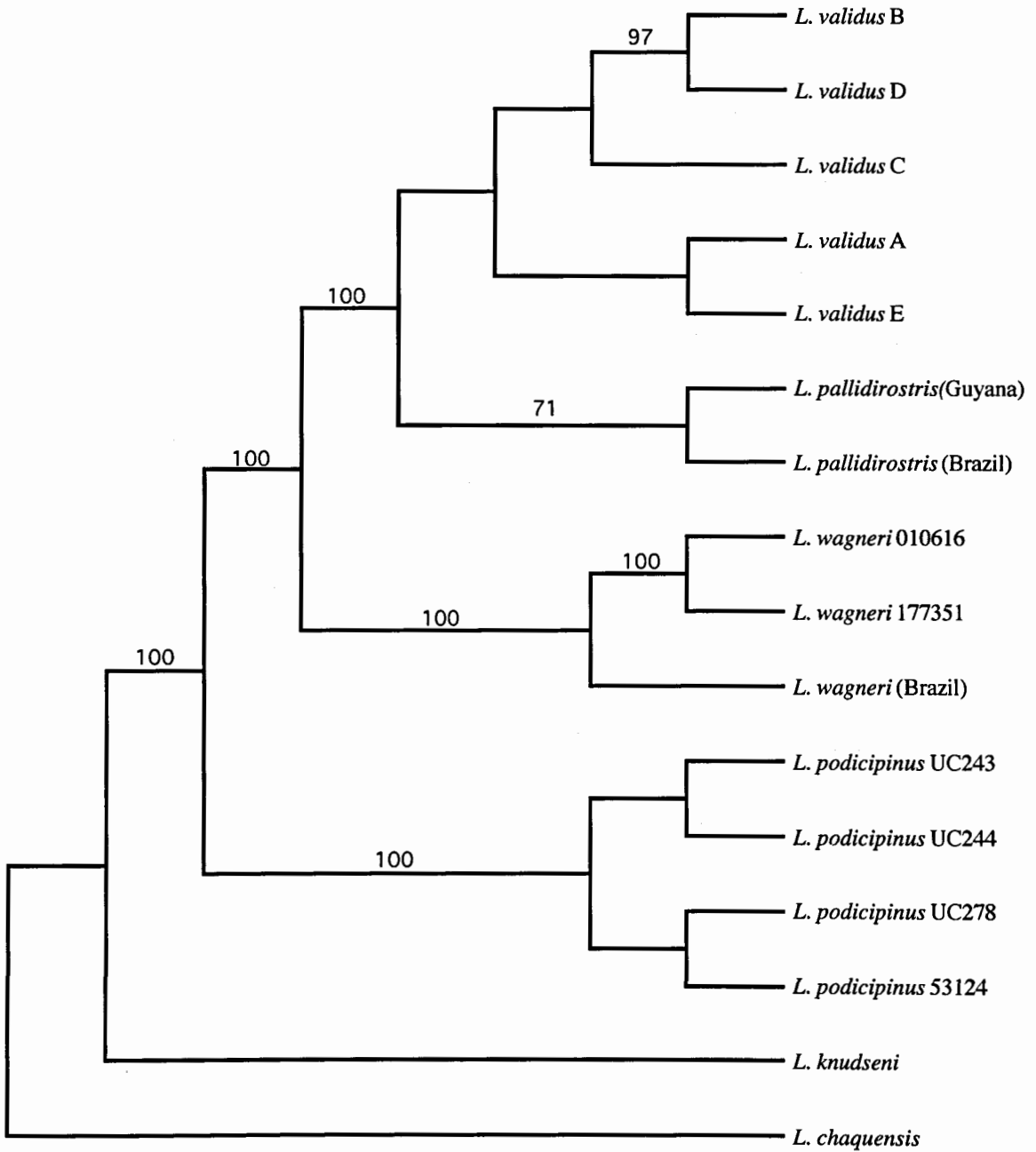




Figure 5. Bayesian consensus tree of 12S gene. Posterior probability values >50% are shown above branches. See text for clade descriptions.



low posterior support.

### 16S rRNA gene

*Sequence divergence.*— Analysis of the 16S sequences grouped the 50 *Leptodactylus validus* samples into seven haplotypes (Table VI). The haplotypes scored showed eleven (0.78%) variable sites among the *L. validus* samples. Haplotype A includes all of the samples from Grenada (n=20) and St. Vincent (n=13). Haplotype B consists of samples from Trinidad (n=11), and haplotype C contains the samples from Tobago (n=2). Haplotypes D–G each consist of a single sample from Trinidad. The aligned 1419 bp fragment of the 16S rRNA gene correspond to positions 3093–4548 in the *Xenopus laevis* genome (Roe et al., 1985; Gen-Bank no.: M10217). A portion of the 16S gene containing multiple indels could not be unambiguously aligned. Subsequently, the 28 bp region was excluded from the analyses (Appendix II). Base frequencies for the ingroup and outgroup taxa are presented in Table VII. Corrected pair-wise genetic distances among *L. validus* samples ranged from 0–0.36%. The two *L. pallidirostris* samples

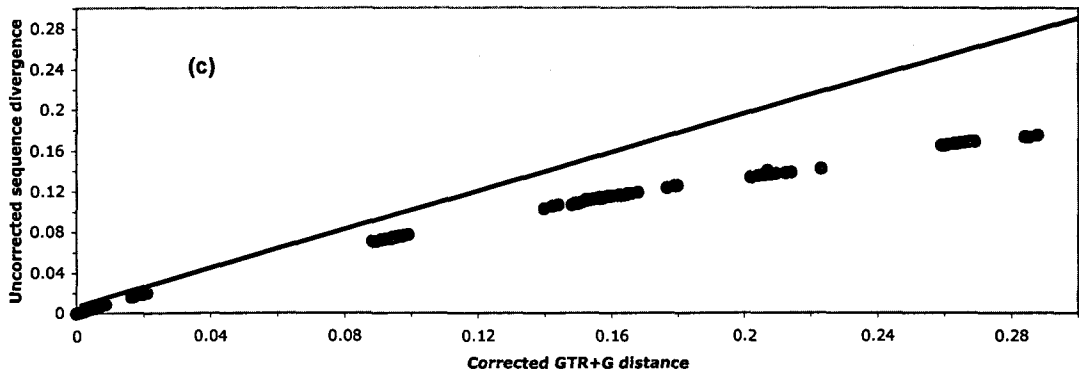
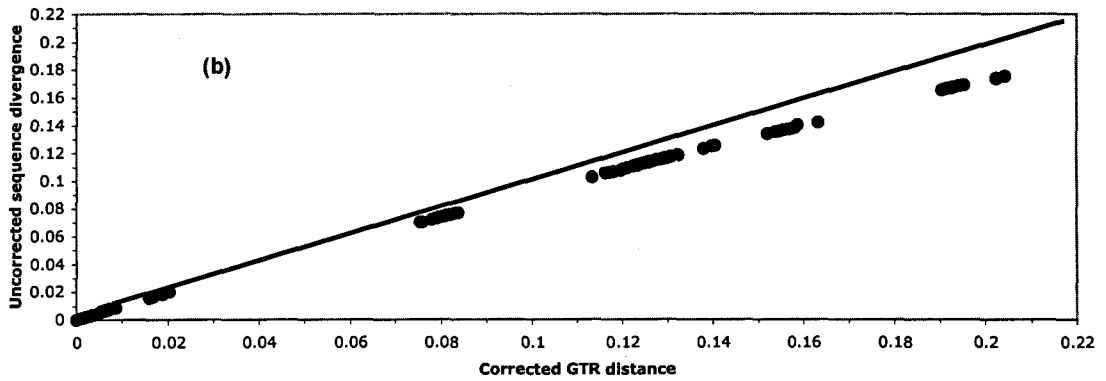
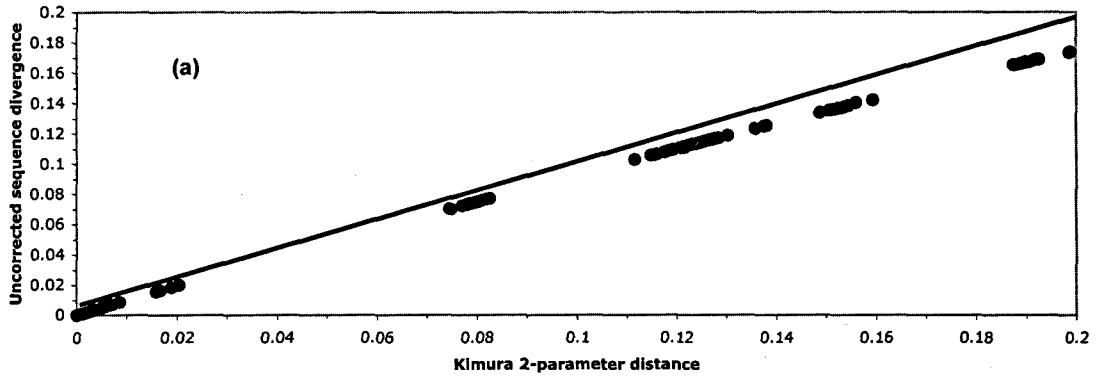
differed by 0.51%. Genetic distances between the *L. pallidirostris* and *L. validus* samples ranged from 0.14–0.73%. Genetic distances among samples of *L. podicipinus* were between 0–2.1% and ranged between 15.22–16.79% when compared with *L.*

*validus/pallidirostris*. Distances among samples of *L. wagneri* were 0.22–1.73% and 8.84–9.92% when compared with *L.*

*validus/pallidirostris*. Distances between *L. validus/pallidirostris* and the outgroups ranged between 20.2–20.96% when compared with *L. chaquensis* and 25.93–26.75% with *L. knudseni*. Corrected pair-wise genetic distances are presented in Table VIII and absolute base differences among samples are provided in Table IX. Plots of pair-wise uncorrected p-distances versus K2p distances were nearly linear, indicating a low degree of transitional saturation (Fig. 6a).

Similarly, the comparison of uncorrected p-distances with corrected GTR divergences did not reveal saturation of base substitutions (Fig. 6b). However, in the comparison of uncorrected p-distances with corrected GTR+ $\Gamma$  distances, the scatter plot points began to depart from the diagonal at higher levels of sequence divergence (Fig. 6c).

Figure 6. Plots depicting relative rates of (a) transitional saturation and (b) substitutional saturation using pair-wise genetic distances corrected under the GTR and (c) GTR+ $\Gamma$  evolutionary models among 16S gene sequences.



*Maximum parsimony analyses.*— Maximum parsimony analysis of the 16S data considering gaps as missing data resulted in a single tree (L=544, CI=0.849) (Fig. 7). Of the 1419 unambiguously aligned bp sequenced, 368 (25.9%) characters were variable and 244 (17.2%) were parsimony-informative. *Leptodactylus validus* and *L. pallidirostris* formed a monophyletic group with 100% bootstrap support. Within this clade, the *L. pallidirostris* sample from Brazil appeared basal to a strongly supported clade that united the *L. pallidirostris* sample from Guyana with a weakly supported *L. validus* subclade. An analysis considering gaps as a fifth character state recovered four minimum-length trees (L=588; CI=0.842; 387 [27.3%] variable characters and 261 [15.2%] parsimony-informative). Topology of the strict consensus tree (Fig. 8) was similar to that of the previous analysis however, the *L. pallidirostris* from Guyana forms an unresolved polytomy with the *L. validus* samples.

*Weighted maximum parsimony analyses.*— An analysis applying a 2:1 (tv:ti) weighting scheme recovered a single tree (L=751; CI=0.856; 368 [25.9%] variable characters and 249 [17.6%]

Figure 7. Consensus tree (50% MR) topology obtained for the 16S data in both unweighted MP analyses, weighted tv:ti and stem-loop MP analyses using gaps as missing data, and weighted ML analysis using the GTR+ $\Gamma$  evolutionary model. Bootstrap values shown above branches correspond to those observed in the unweighted MP analysis considering gaps as missing data. See text for clade descriptions.



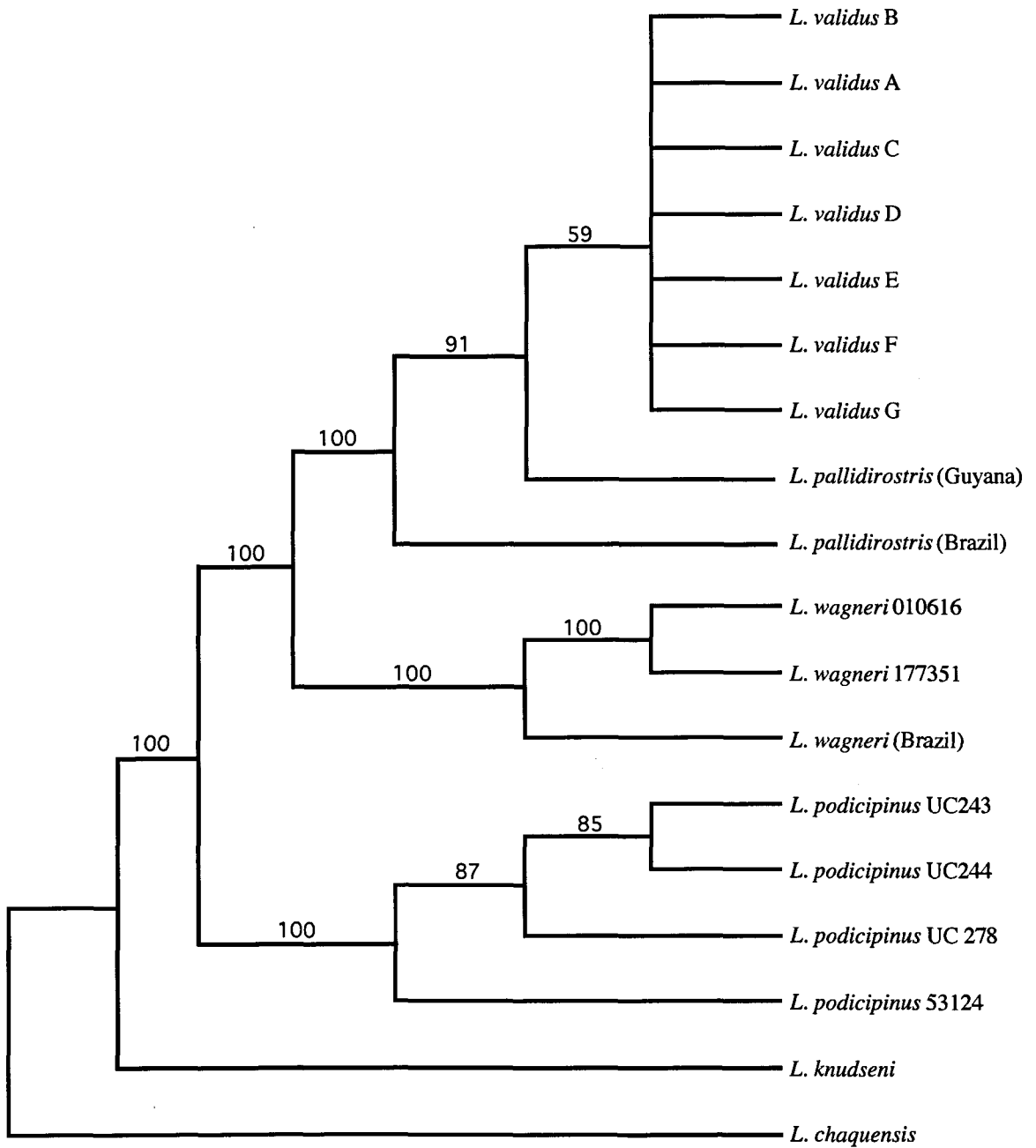
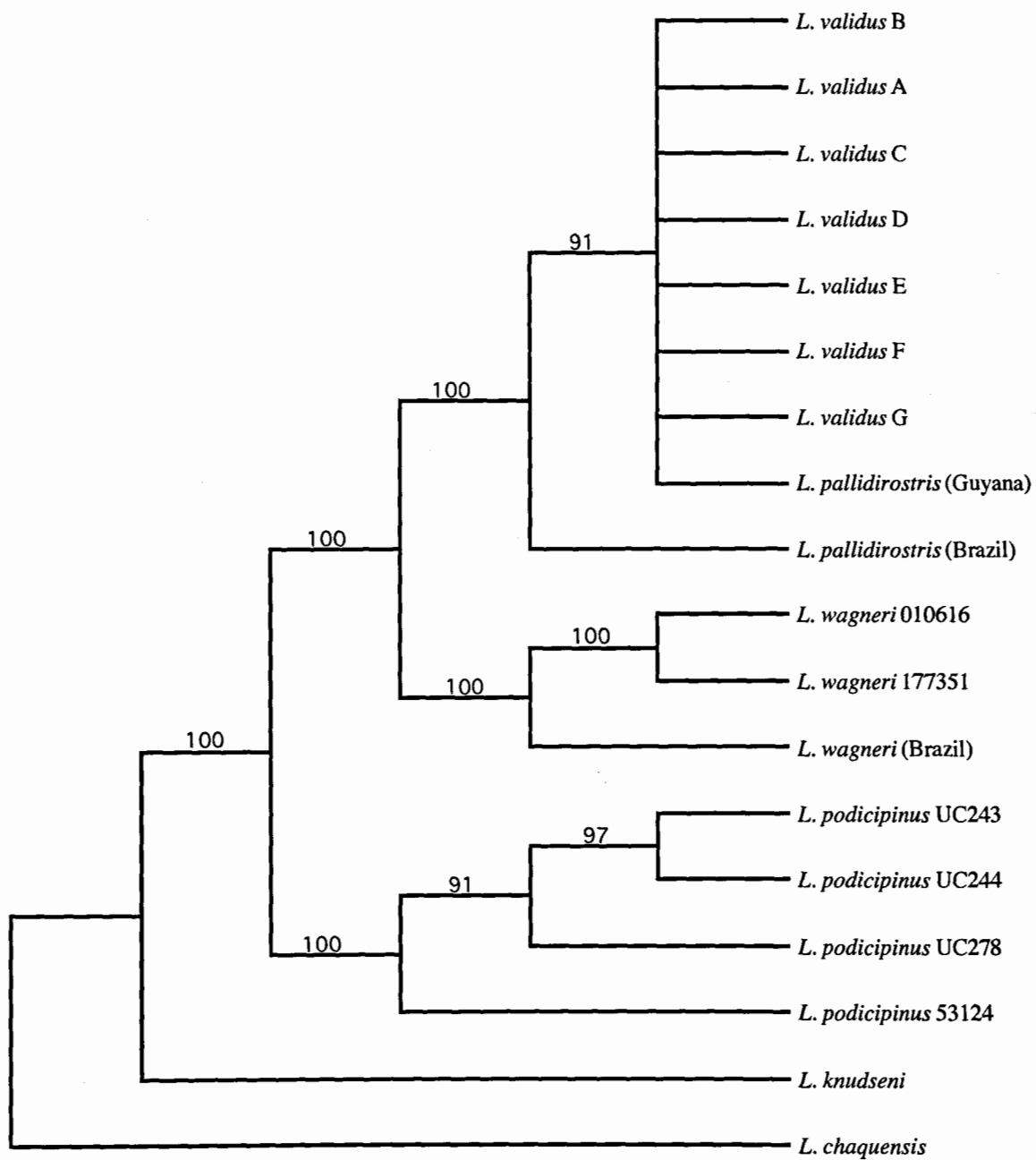


Figure 8. Consensus tree (50% MR) topology obtained for the 16S data in unweighted MP analysis considering gaps as a fifth character state and the weighted ML analysis performed under the TrN+ $\Gamma$  evolutionary model. Bootstrap values shown above branches correspond to those observed in the unweighted MP analysis considering gaps as a fifth character state. See text for clade descriptions.



parsimony-informative). Weighting loop positions (n=755) to stem positions (n=664) and considering gaps as missing data also resulted in a single tree (L=917; CI=0.855; 368 [25.9%] variable characters and 244 [17.2%] parsimony-informative). Topologies of the trees from these analyses were identical to the one presented in Fig. 7. Using gaps as a fifth character state under the same weighting scheme, a single tree was obtained (L=1004; CI=0.847; 387 [27.3%] variable characters and 261 [18.4%] parsimony-informative). The strict consensus tree was similar to the one depicted in Fig. 8 with the addition of a subclade formed by the *L. pallidirostris* sample from Guyana and two *L. validus* samples from Trinidad (Fig. 9).

*Maximum likelihood analyses.*— Maximum likelihood analysis under the TrN+G model parameters resulted in a bootstrap 50% MR consensus tree topology similar to the one depicted in Fig. 7, with less resolution within the *L. podicipinus* clade (Fig. 10). A similar tree was obtained under the GTR+ $\Gamma$  model parameters, with relationships among samples within the *L. podicipinus* clade differing slightly (Fig. 11). ML analysis considering secondary

Figure 9. Weighted MP tree obtained for the 16S data using gaps as a fifth character state. Bootstrap values are shown above branches. See text for clade descriptions.

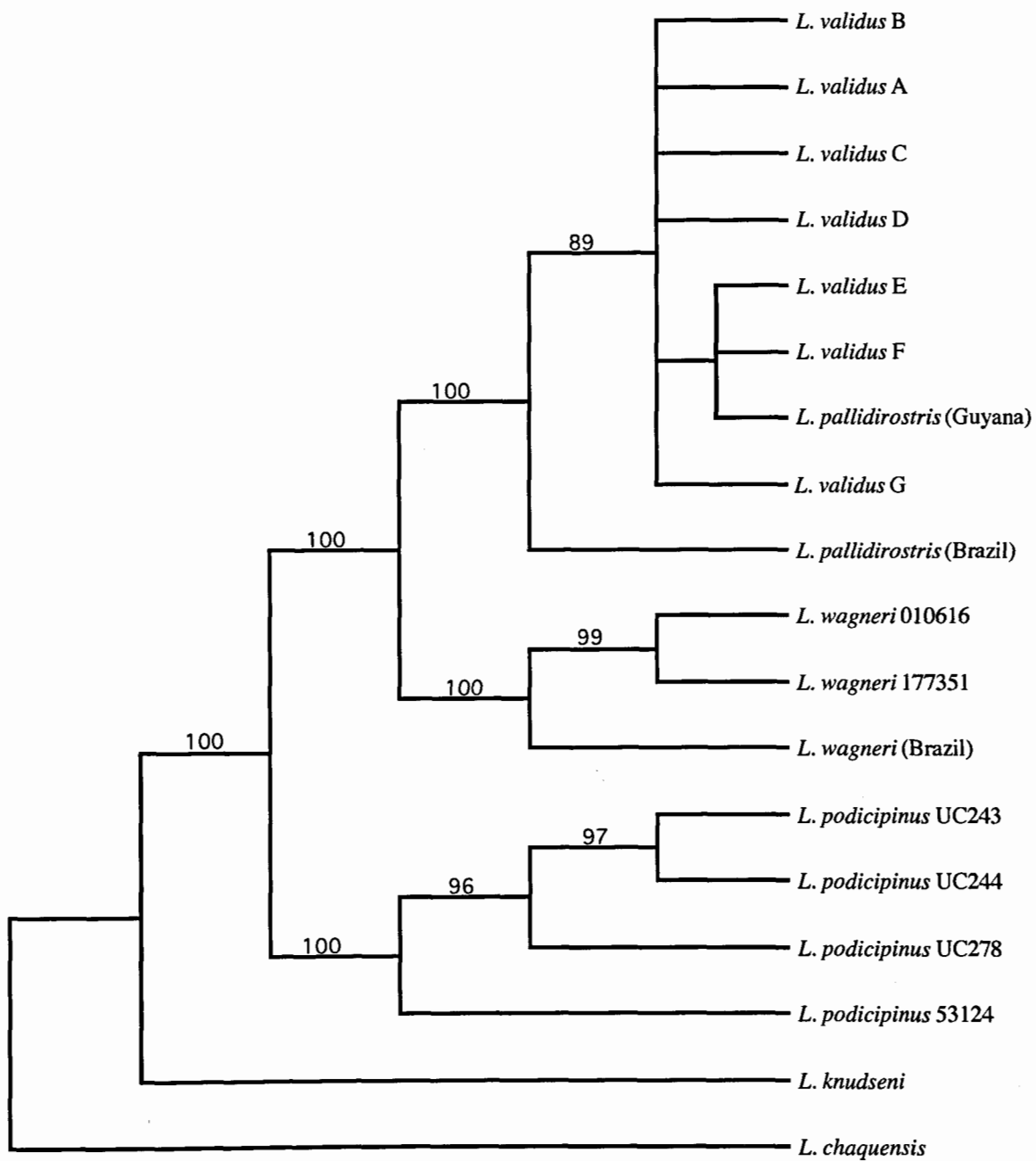


Figure 10. Unweighted ML tree obtained for the 16S data using the TrN+ $\Gamma$  evolutionary model. Bootstrap values are shown above branches. See text for clade descriptions.

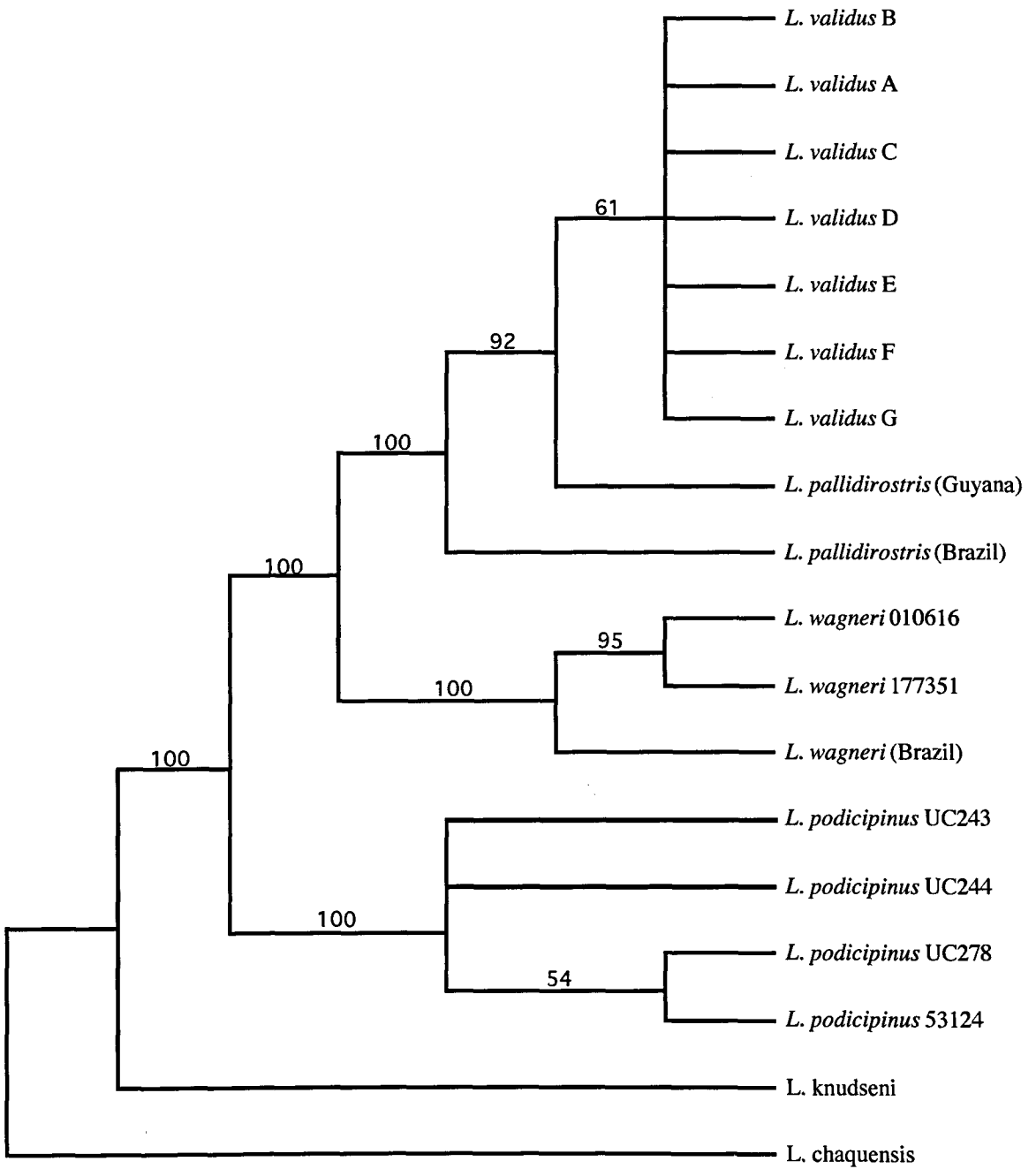
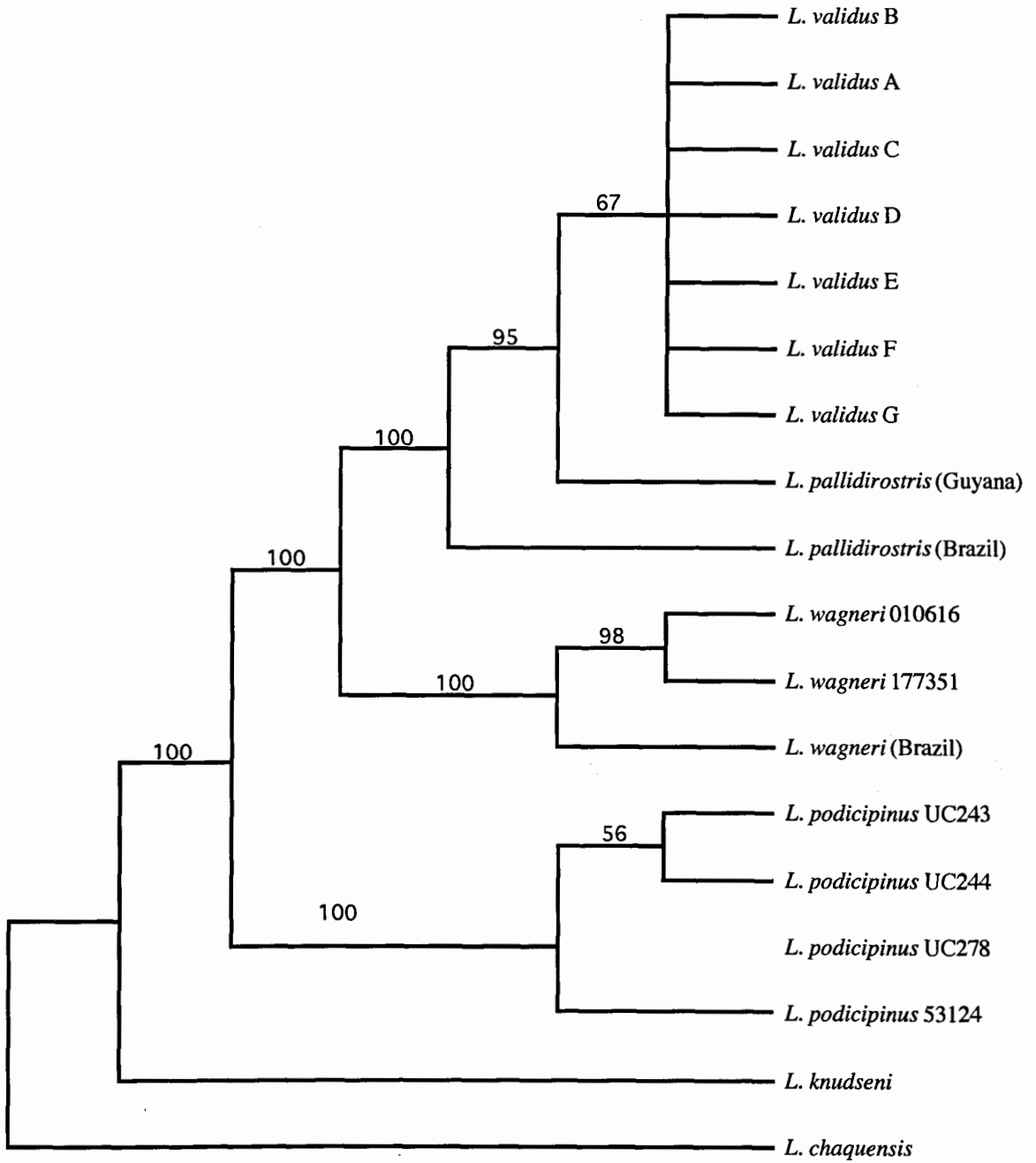




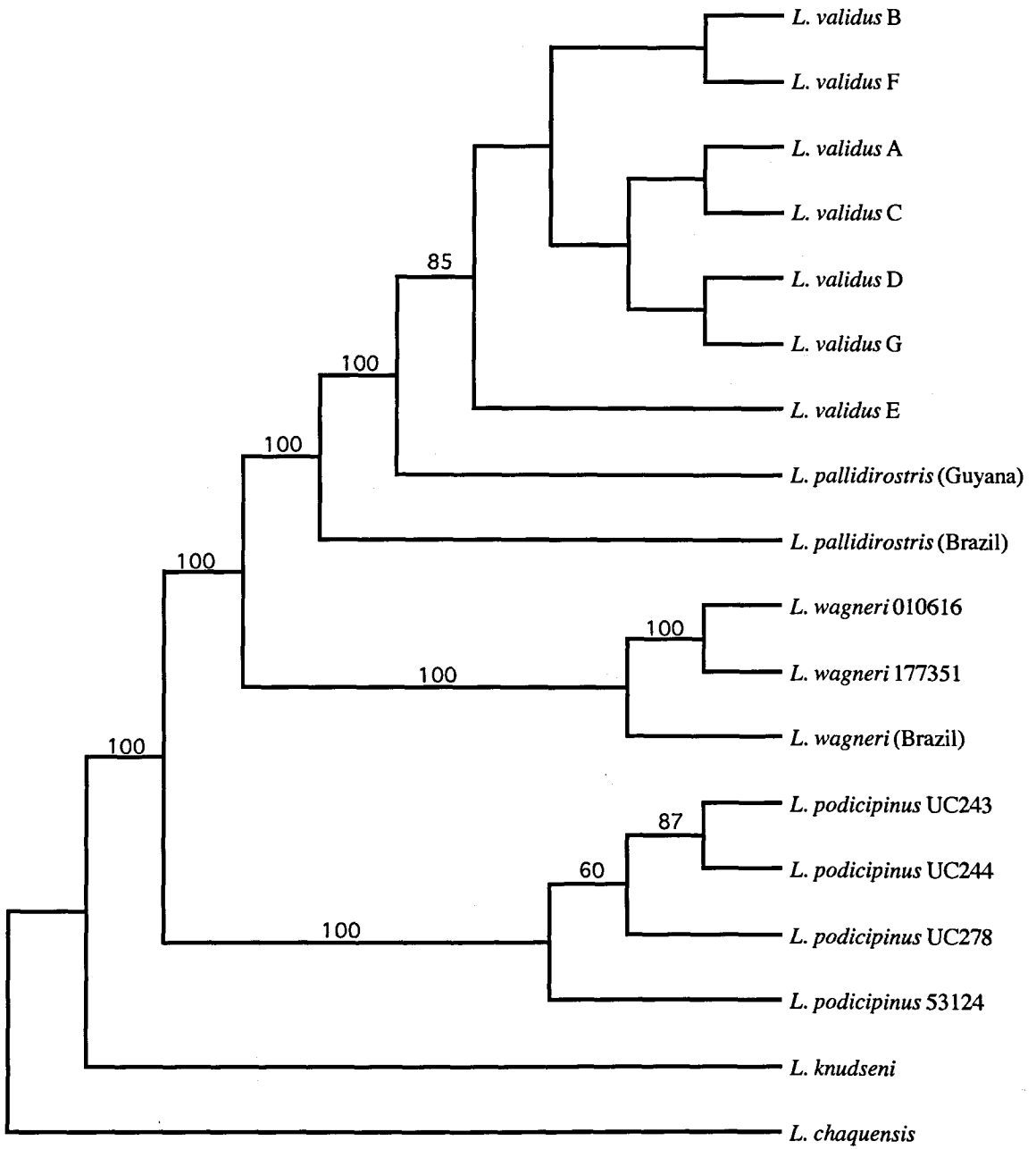
Figure 11. Unweighted ML tree obtained for the 16S data using the GTR+ $\Gamma$  evolutionary model. Bootstrap values are shown above branches. See text for clade descriptions.



structure and weighting loop:stem positions (2:1) under the TrN+G model parameters, resulted in a bootstrap 50% MR consensus tree with topology identical to the tree observed in Fig. 8, with less support for the relationships depicted among the *L. podicipinus* samples. The topology of the bootstrap 50% MR consensus tree obtained in the weighted ML analysis under the GTR+ $\Gamma$  model parameters was identical to the one presented in Fig. 7.

*Bayesian analysis.*— A Bayesian analysis of the 16S dataset was performed using the GTR+ $\Gamma$  model setting. Convergence of the Log likelihood values among the seven MCMC chains occurred within 16,000 generations of sampling, consequently the burn-in value was set to discard the first 1,600 trees sampled. The consensus tree obtained by MrBayes (Fig. 12) was similar to the bootstrap strict consensus trees described in the MP analyses when gaps were considered as missing data. There was 100% posterior support for a monophyletic *L. pallidirostris*/*L. validus* clade. The *L. pallidirostris* sample from Brazil appeared basal to a subclade in which a dichotomy was formed between the *L. pallidirostris* sample from Guyana and *L. validus*. There was 100% posterior support for this

Figure 12. Bayesian consensus tree of 16S gene. Posterior probability values >50% are shown above branches. See text for clade descriptions.



subclade, however support for the relationships among the *L. validus* samples was extremely low.

#### Combined data

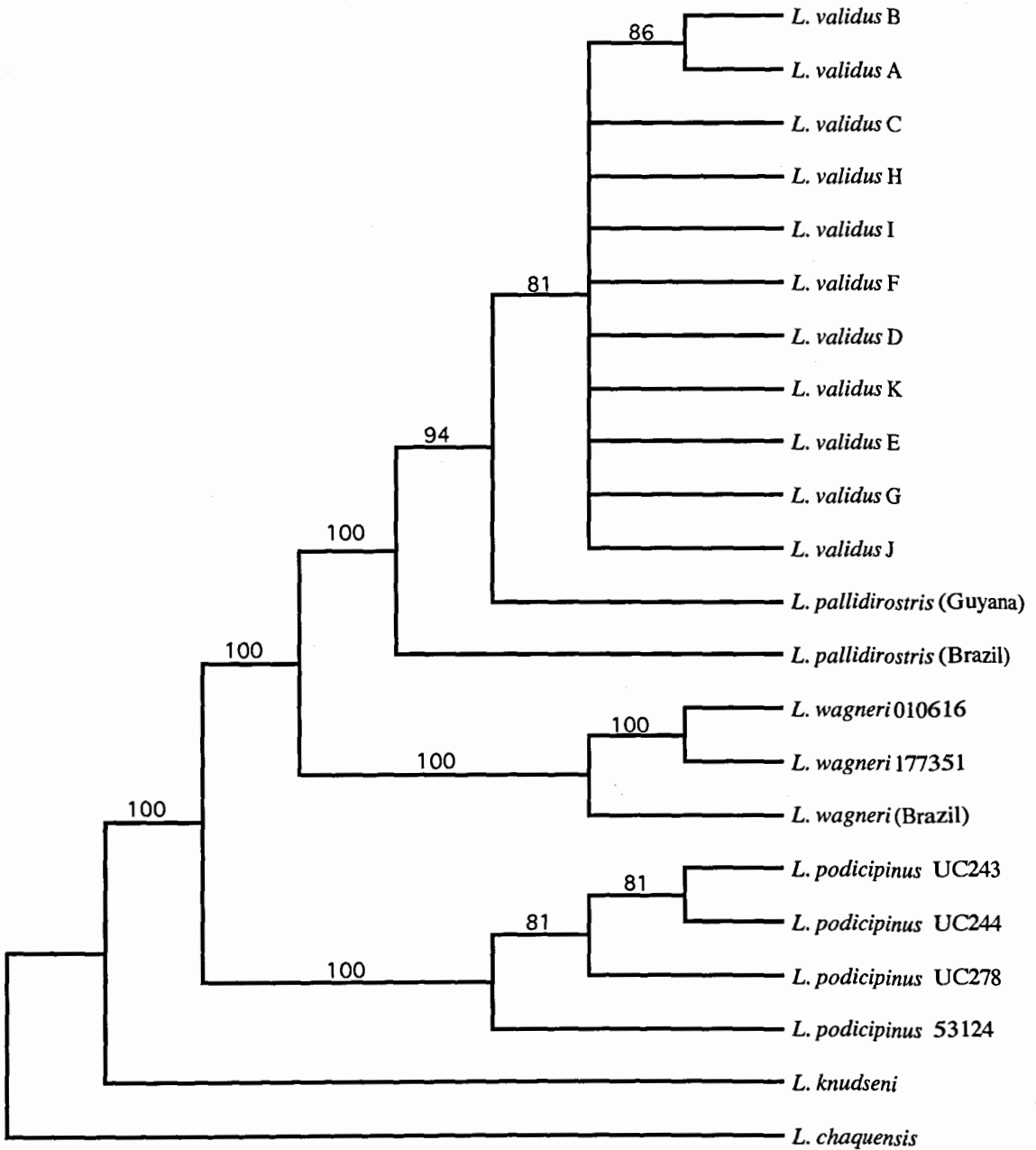
*Sequence divergence.*— Analysis of the combined sequence data grouped the 50 *Leptodactylus validus* samples into eleven haplotypes (Table X). The haplotypes scored showed sixteen (0.71%) variable sites among the *L. validus* samples. Haplotype A includes samples from Grenada (n=9) and St. Vincent (n=8). Haplotype B consists of samples from Grenada (n=11) and St. Vincent (n=5). Haplotype C includes samples from Trinidad (n=8), and haplotype D contains the samples from Tobago (n=2). Haplotypes E–K each consist of a single sample from Trinidad. Corrected pair-wise genetic distances among *L. validus* samples ranged from 0–0.32%. The two *L. pallidirostris* samples differed by 0.41%. Genetic distances between the *L. pallidirostris* and *L. validus* samples ranged from 0.14–0.64%. Genetic distances among samples of *L. podicipinus* were between 0.05–1.64% and ranged between 13.45–14.54% when compared with *L. validus/pallidirostris*.

Distances among samples of *L. wagneri* were 0.23–1.65% and 7.41–8.04% between *L. validus/pallidirostris*. Distances between *L. validus/pallidirostris* and the outgroups ranged between 16.93–17.55% compared with *L. chaquensis* and 22.27–22.83% with *L. knudseni*. Corrected pair-wise genetic distances are presented in Table XI and absolute base differences among samples are provided in Table XII.

*Maximum parsimony analyses.*— Maximum parsimony analysis of the Combined data considering gaps as missing data resulted in three equally parsimonious trees, (L=765; CI=0.85). Of the 2247 aligned characters sequenced, 527 (23.5%) characters were variable, and 349 (15.5%) were parsimony-informative. The strict consensus tree (Fig. 13) depicts the *L. pallidirostris* and *L. validus* samples as a monophyletic group with 100% support. Within this clade, the *L. pallidirostris* sample from Brazil appeared basal to a strongly supported clade that united the *L. pallidirostris* sample from Guyana with a strongly supported *L. validus* subclade. There is strong support for a subclade formed by the two Grenada/St. Vincent haplotypes within the *L. validus* clade.

Figure 13. Consensus tree (50% MR) topology obtained in all MP analyses of the combined dataset. Bootstrap values shown above branches correspond to those observed in the MP analysis considering gaps as missing data. See text for clade descriptions.





Relationships among the other *L. validus* samples are unresolved.

An analysis considering gaps as a fifth character state recovered three minimum-length trees (L=815; CI=0.84; 549 [24.4%] variable characters and 370 [16.5%] parsimony-informative). Topology of the strict consensus tree was identical to the one presented in Fig. 13.

*Weighted maximum parsimony analyses.*— An analysis applying a 2:1 (tv:ti) weighting scheme recovered three minimum-length trees (L=1041; CI=0.856; 527 [23.5%] variable characters and 355 [15.8%] parsimony-informative). Weighting loop positions (n=1135) to stem positions (n=1112), and considering gaps as missing data, resulted in three minimum-length trees (L=1269; CI=0.850; 527 [23.5%] variable characters and 349 [15.5%] parsimony-informative). Topologies of the strict consensus trees obtained in these analyses were identical to the one presented in Fig. 13. Using gaps as a fifth character state under the same weighting scheme, a single tree was obtained (L=1379; CI=0.839; 549 [24.4%] variable characters and 370 [16.5%] parsimony-

informative. Topology of this tree, as well as the two weighted strict consensus trees were identical to those obtained in the unweighted analyses Fig. 13.

*Maximum likelihood analyses.* — Maximum likelihood analysis of the combined data implemented under the TrN+G model parameters resulted in a bootstrap 50% MR consensus tree topology similar to the one obtained in the MP analyses, with relationships among samples within the *L. podicipinus* clade differing slightly (Fig. 14). A similar tree was obtained using the GTR+I+ $\Gamma$  model parameters, with better resolution demonstrated among samples within the *L. podicipinus* clade (Fig. 15). ML analyses considering secondary structure and weighting loop:stem positions (2:1), using both evolutionary models resulted in bootstrap 50% MR consensus trees identical to the tree presented in Fig. 14.

*Bayesian analysis.*— A Bayesian analysis was conducted under the GTR+I+ $\Gamma$  model setting. Convergence of the Log likelihood values among the seven MCMC chains occurred within 40,000 generations of sampling, consequently the burn-in value was set to discard the first 4,000 trees sampled. The consensus tree obtained

Figure 14. ML Tree topology obtained for the combined dataset from unweighted analysis using the TrN+ $\Gamma$  evolutionary model and weighted analyses using both models. Bootstrap values shown above branches correspond to those observed in the unweighted analysis using the TrN+ $\Gamma$  model. See text for clade descriptions.

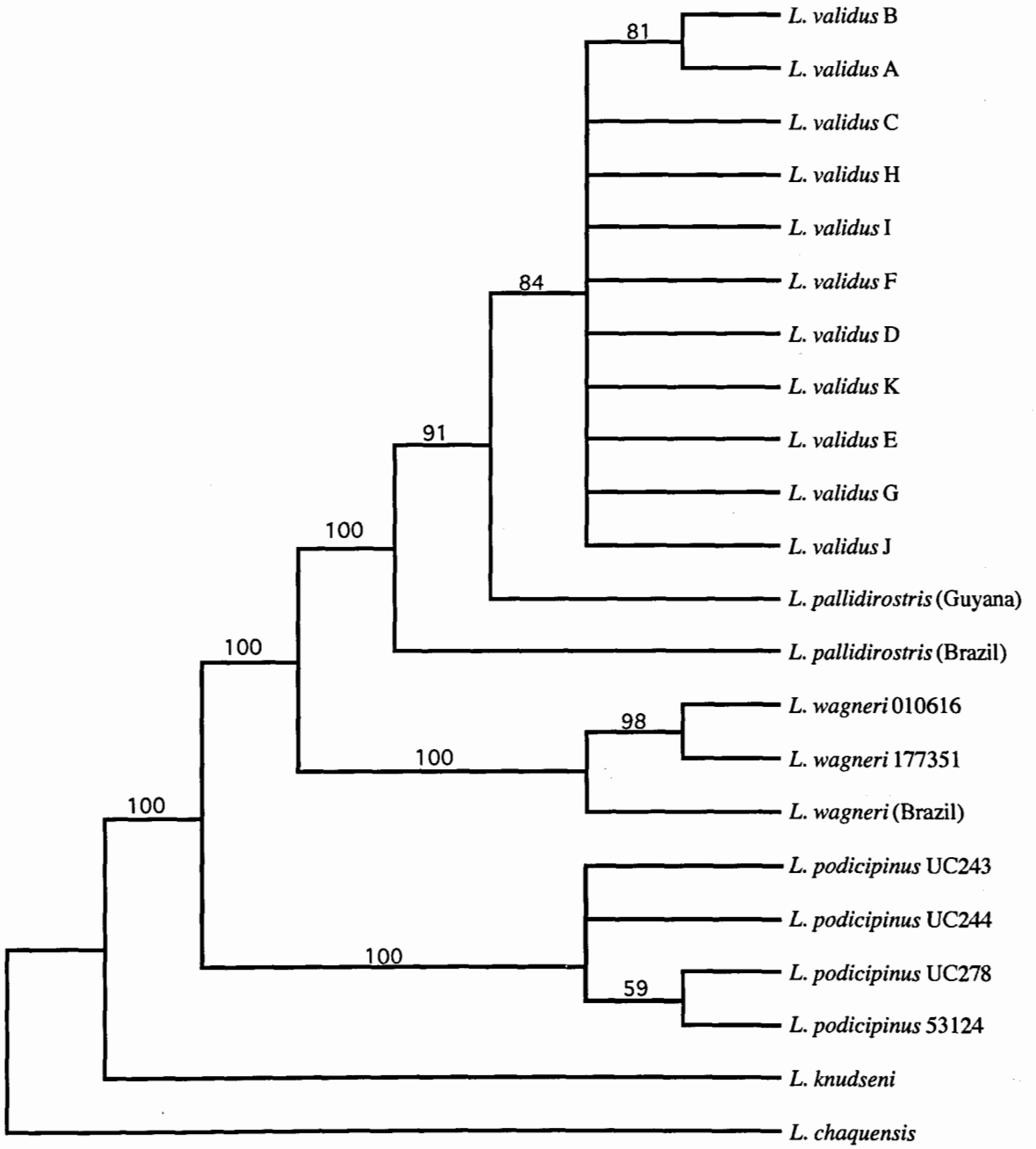
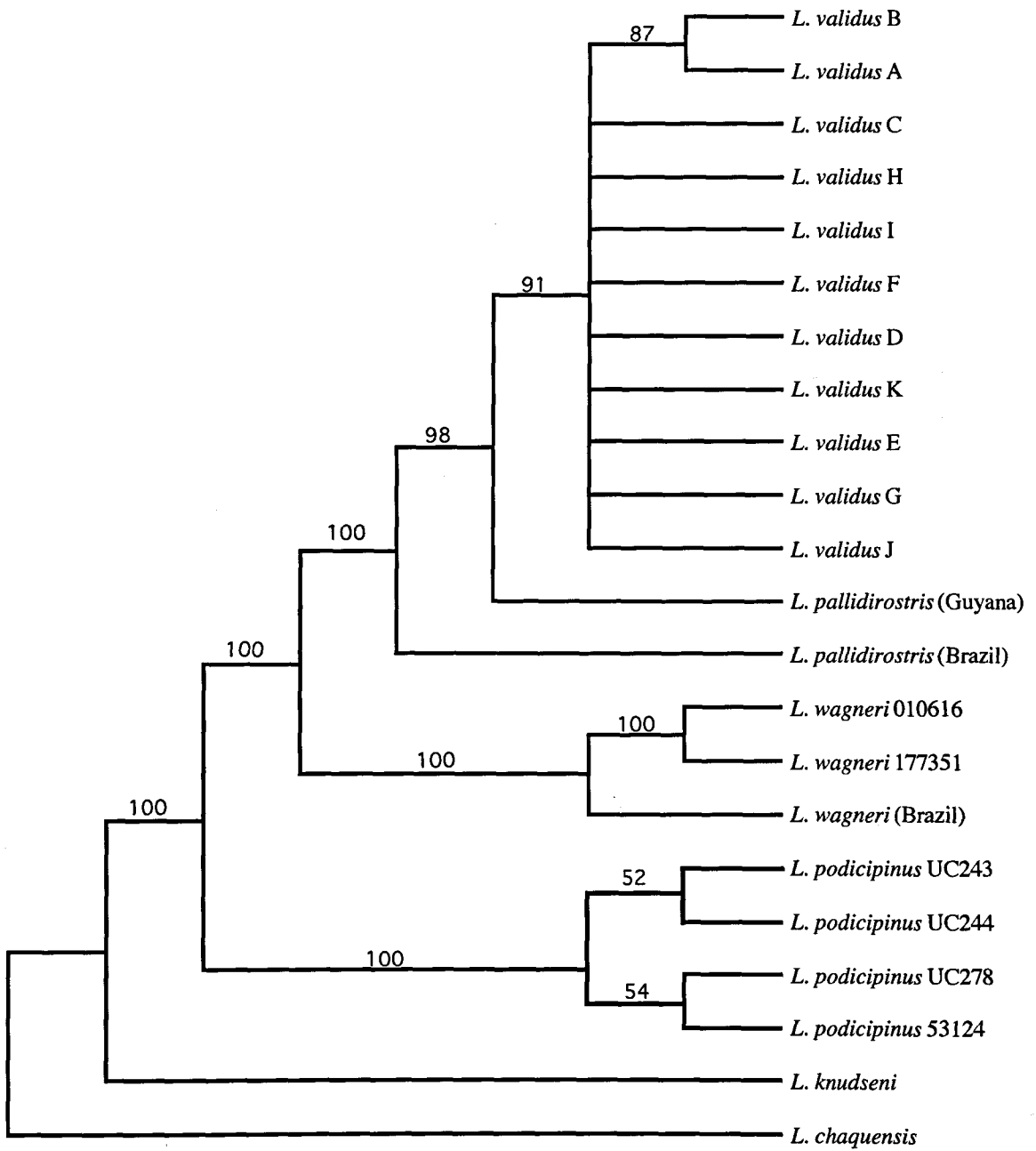


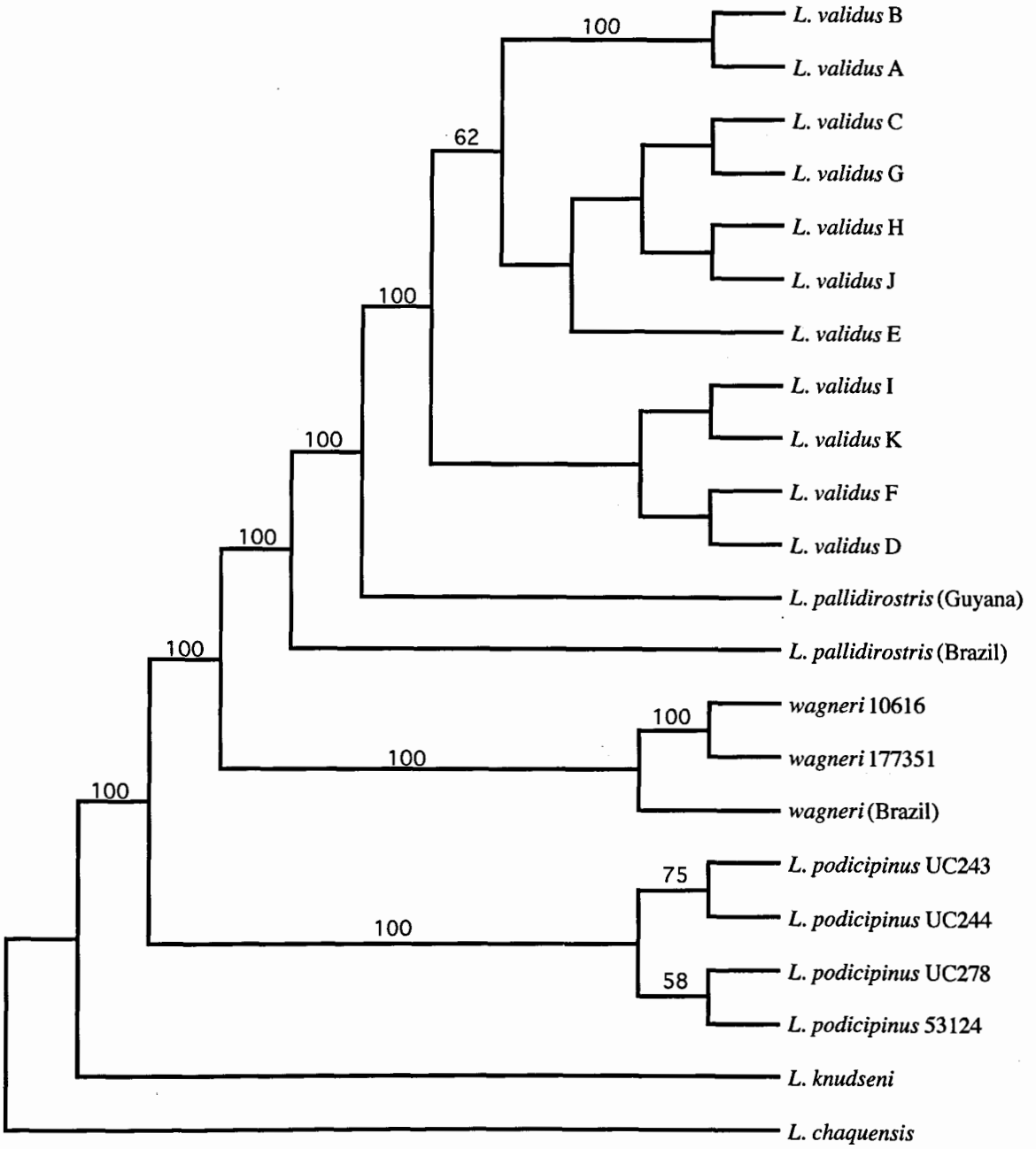
Figure 15. Unweighted ML tree obtained for the combined dataset using the GTR+I+ $\Gamma$  evolutionary model. Bootstrap values are shown above branches. See text for clade descriptions.



by MrBayes (Fig. 16) was similar to the unweighted ML tree obtained using the GTR+I+ $\Gamma$  model parameters, with better resolution among the *L. validus* samples. With the exception of the subclade consisting of the Grenada/St. Vincent haplotypes however, there was little posterior support for any of the relationships depicted among the samples within the *L. validus* subclade.



Figure 16. Bayesian consensus tree of combined dataset. Posterior probability values >50% are shown above branches. See text for clade descriptions.



## Discussion

Using morphological characters and call data, Heyer (1994) assessed variation among species within the *Leptodactylus podicipinus-wagneri* complex. However, morphological data were insufficient to resolve some questions involving the species boundaries in this complex. Consequently, the status of *L. pallidirostris* and *L. validus* within this complex remained a major unresolved problem (Heyer, 1994). Furthermore, the presence of *Leptodactylus validus* on the continental islands of Trinidad and Tobago and the oceanic islands of the Lesser Antilles, along with its absence from the mainland of South America, is inconsistent with the distribution patterns normally observed for other fauna of this region. Heyer (1994) described four possible distribution patterns expected for a species in this region: 1) occurring on the mainland of South America, Trinidad, Tobago, and the Lesser Antilles, 2) present in mainland South America, Trinidad, and Tobago but not on the Lesser Antilles, 3) found only in Trinidad and Tobago, or 4) found only in the Lesser Antilles. He suggested that either *L. validus* was present on the South American mainland or alternatively, that

*L. validus* might represent two or more closely related, morphologically similar species throughout its distribution.

Molecular techniques involving slowly evolving mitochondrial rRNA genes (12S and 16S), have frequently been employed in investigations of the phylogenetic relationships and biogeography among closely related taxa (Hass et al., 1993; Pramuk et al., 2001; Fritsch, 2003; Hower and Hedges, 2003). The use of these genes has also proven useful in revealing the presence sibling species among morphologically indistinguishable species (Richards et al., 2000; Glaw and Vences 2002).

All analyses performed in this study strongly supported a monophyletic group consisting of the *L. pallidirostris* and *L. validus* samples. The combined analyses revealed some genetic structuring among populations within this clade, i.e., genetically distant samples were also geographically distant. For example, a close relationship was demonstrated among samples of *L. validus* from the Lesser Antilles. Likewise, the *L. pallidirostris* sample from Guyana appeared more closely related to the *L. validus* samples than the *L. pallidirostris* sample from Brazil. Comparing corrected pair-wise

genetic distances between the *L. validus* samples and the *L. pallidirostris* samples from each dataset resulted in the following genetic distances: <0.37% different for the *L. pallidirostris* sample from Guyana and <0.73% different for the *L. pallidirostris* sample from Brazil. Less than 0.5% sequence divergence was observed among the *L. validus* samples. These low levels of sequence divergence suggest *L. pallidirostris* and *L. validus* to be conspecific, supporting the hypothesis describing *L. validus* as a single species, distributed throughout the Lesser Antilles, Trinidad, Tobago, and mainland South America.

Advertisement call is another characteristic commonly analyzed in anuran systematic studies. These calls possess features that are species-specific in *Leptodactylus*, and in anurans in general, and are often a reliable indicator of species boundaries in frogs (Heyer and Straughan, 1976; Heyer, 1978, 1979, 1994; Heyer et al., 1996; Wieczorek and Channing, 1997; Hertwig et al., 2004; Camargo et al., submitted; de Sá et al., submitted, Heyer et al., submitted). The call data analyzed from Venezuelan populations of *L. pallidirostris* were similar to the advertisement call of *L. validus*

(Heyer, 1994), providing additional support for the conspecific status of the two species. Heyer indicated that a conservative approach was taken with regard to *L. pallidirostris*, and there was enough morphological variation present to suggest that more than one species may be present among the samples he examined.

Additionally, Heyer (1994) explained that some of the Venezuelan samples he examined and assigned to *L. pallidirostris* closely resembled Trinidad and Tobago samples that he designated as *L. validus*, suggesting that the two populations may be conspecific.

*Leptodactylus pallidirostris* was first described from Katarbo, Guyana, by Lutz in 1930, based on coloration and pattern, however Lutz repeatedly referred to the species' resemblance to *L. validus*. This is in agreement with the present study in which *L. pallidirostris* and *L. validus* appears to be a single taxon. *Leptodactylus validus* was described by Garman in 1877, therefore this name has scientific priority over *L. pallidirostris*. Consequently, *L. pallidirostris* is placed in the synonymy of *L. validus*.

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Table I. Primers used for amplification (1) and sequencing (2) of 12S and 16S rDNA fragments.

Name	Sequence (5'-3')	
12SA-L <sup>1,2</sup>	AAACTGGGATTAGATACCCCACTAT	(Palumbi et al., 1991)
12SB-H <sup>1,2</sup>	GAGGGTGACGGGCGGTGTGT	(Palumbi et al., 1991)
12Stp <sup>hef</sup> <sup>1,2</sup>	ATAGC (A/G)CTGAA (A/G)A (C/T)GCT (A/G)AGATG	
12SRds1 <sup>1,2</sup>	GGTACCGTCAAGTCCTTTGGGTT	
12L13 <sup>1,2</sup>	TTAGAAGAGGCAAGTCGTAACATGGTA	(Feller & Hedges 1998)
Hedges16H10 <sup>1,2</sup>	TGATTACGCTACCTTTGCACGGT	(Hedges 1994)
16Sar-L <sup>1,2</sup>	CGCCTGTTTACCAAAAACAT	(Palumbi et al., 1991; Graybeal 1997)
16Sbr-H <sup>2</sup>	CCGGTCTGAACTCAGATCACGT	(Palumbi et al., 1991)
16Sd <sup>1</sup> (Hedges16H1)	CTCCGGTCTGAACTCAGATCACGTAG	(Hedges, 1994)

Table II. 12S rDNA sequence haplotypes for *L. validus* samples from: Grenada (Gren), St. Vincent (StVn), Tobago (Tobo), and Trinidad (Trin). Haplotypes A and B represent multiple samples. Asterisks indicate samples used to represent a haplotype. Haplotypes C–E each consist of a single sample: (C) Trin175424, (D) Trin196728, and (E) Trin196886.

A	B
Gren006881	Gren196977
Gren006882	Gren196979
Gren006883	Gren196980
Gren006939	Gren197000
Gren196978	Gren197001
Gren196999	Gren197002
Gren197005	Gren197003
Gren197006	Gren197004
Gren197017	Gren197007
StVn056421	Gren197008
StVn056490	Gren197044
StVn056561	StVn196894
StVn056562	StVn196896
StVn056612	StVn196897
*StVn056613	StVn196899
StVn196895	StVn196900
StVn196898	Trin175410
Tobo186596	Trin175620
Tobo186597	*Trin196726
Trin196729	Trin196727
	Trin196730
	Trin196731
	Trin196732
	Trin196733
	Trin196734
	Trin196735
	Trin196888

Table III. Base composition percentages of 12S mtDNA gene sequences (averages among samples from each taxa).

	<i>validus</i>	<i>pallidirostris</i>	<i>podicipinus</i>	<i>wagneri</i>	<i>chaquensis</i>	<i>knudseni</i>	All taxa
A	0.31089	0.31280	0.22970	0.31316	0.31356	0.32121	0.31094
T	0.23857	0.23551	0.16515	0.22720	0.23123	0.21818	0.23625
G	0.20413	0.20411	0.15909	0.20581	0.20339	0.20242	0.20473
C	0.24640	0.24759	0.19606	0.25384	0.25182	0.25818	0.24807

Table IV. Pair-wise genetic distances for the 12S dataset. Distances corrected using the general time-reversible model with gamma distributed rates for variable sites.

	1	2	3	4	5	6	7	8
1 Gren006881	-							
2 Gren006882	0.00000	-						
3 Gren006883	0.00000	0.00000	-					
4 Gren006939	0.00000	0.00000	0.00000	-				
5 Gren196977	0.00122	0.00122	0.00122	0.00122	-			
6 Gren196978	0.00000	0.00000	0.00000	0.00000	0.00122	-		
7 Gren196979	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	-	
8 Gren196980	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	-
9 Gren196999	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
10 Gren197000	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
11 Gren197001	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
12 Gren197002	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
13 Gren197003	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
14 Gren197004	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
15 Gren197005	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
16 Gren197006	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
17 Gren197007	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
18 Gren197008	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
19 Gren197017	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
20 Gren197044	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
21 StVn056421	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
22 StVn056490	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
23 StVn056561	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
24 StVn056562	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
25 StVn056612	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
26 StVn196895	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
27 StVn196896	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
28 StVn196897	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
29 StVn196898	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
30 StVn196899	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
31 StVn196900	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
32 Tobo186597	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
33 Trin196726	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
34 Trin196727	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
35 Trin196728	0.00244	0.00244	0.00244	0.00244	0.00122	0.00244	0.00122	0.00122
36 Trin196729	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
37 Trin196730	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
38 Trin196731	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
39 Trin196732	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
40 Trin196733	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
41 Trin196734	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
42 Trin196888	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
43 StVn056613	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
44 StVn196894	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
45 Trin175424	0.00243	0.00243	0.00243	0.00243	0.00366	0.00243	0.00366	0.00366
46 Tobo186596	0.00000	0.00000	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122
47 Trin196886	0.00122	0.00122	0.00122	0.00122	0.00244	0.00122	0.00244	0.00244
48 Trin175410	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
49 Trin175620	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
50 Trin196735	0.00122	0.00122	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000
51 pall207682	0.00122	0.00122	0.00122	0.00122	0.00245	0.00122	0.00245	0.00245
52 pallWR8626	0.00368	0.00368	0.00368	0.00368	0.00492	0.00368	0.00492	0.00492
53 wagn010616	0.05218	0.05218	0.05218	0.05218	0.05366	0.05218	0.05366	0.05366
54 wagn012969	0.05361	0.05361	0.05361	0.05361	0.05509	0.05361	0.05509	0.05509
55 wagn177351	0.05042	0.05042	0.05042	0.05042	0.05189	0.05042	0.05189	0.05189
56 podicUC243	0.10749	0.10749	0.10749	0.10749	0.10930	0.10749	0.10930	0.10930
57 podicUC244	0.10586	0.10586	0.10586	0.10586	0.10767	0.10586	0.10767	0.10767
58 podicUC278	0.10913	0.10913	0.10913	0.10913	0.11095	0.10913	0.11095	0.11095
59 podic53124	0.10849	0.10849	0.10849	0.10849	0.11032	0.10849	0.11032	0.11032
60 chaq186524	0.12481	0.12481	0.12481	0.12481	0.12550	0.12481	0.12550	0.12550
61 knuds13244	0.17003	0.17003	0.17003	0.17003	0.17219	0.17003	0.17219	0.17219

	9	10	11	12	13	14	15	16
9 Gren196999	-							
10 Gren197000	0.00122	-						
11 Gren197001	0.00122	0.00000	-					
12 Gren197002	0.00122	0.00000	0.00000	-				
13 Gren197003	0.00122	0.00000	0.00000	0.00000	-			
14 Gren197004	0.00122	0.00000	0.00000	0.00000	0.00000	-		
15 Gren197005	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	-	
16 Gren197006	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	0.00000	-
17 Gren197007	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
18 Gren197008	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
19 Gren197017	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	0.00000	0.00000
20 Gren197044	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
21 StVn056421	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	0.00000	0.00000
22 StVn056490	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	0.00000	0.00000
23 StVn056561	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	0.00000	0.00000
24 StVn056562	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	0.00000	0.00000
25 StVn056612	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	0.00000	0.00000
26 StVn196895	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	0.00000	0.00000
27 StVn196896	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
28 StVn196897	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
29 StVn196898	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	0.00000	0.00000
30 StVn196899	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
31 StVn196900	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
32 Tobo186597	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	0.00000	0.00000
33 Trin196726	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
34 Trin196727	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
35 Trin196728	0.00244	0.00122	0.00122	0.00122	0.00122	0.00122	0.00244	0.00244
36 Trin196729	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	0.00000	0.00000
37 Trin196730	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
38 Trin196731	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
39 Trin196732	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
40 Trin196733	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
41 Trin196734	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
42 Trin196888	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
43 StVn056613	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	0.00000	0.00000
44 StVn196894	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
45 Trin175424	0.00243	0.00366	0.00366	0.00366	0.00366	0.00366	0.00243	0.00243
46 Tobo186596	0.00000	0.00122	0.00122	0.00122	0.00122	0.00122	0.00000	0.00000
47 Trin196886	0.00122	0.00244	0.00244	0.00244	0.00244	0.00244	0.00122	0.00122
48 Trin175410	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
49 Trin175620	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
50 Trin196735	0.00122	0.00000	0.00000	0.00000	0.00000	0.00000	0.00122	0.00122
51 pall207682	0.00122	0.00245	0.00245	0.00245	0.00245	0.00245	0.00122	0.00122
52 pallWR8626	0.00368	0.00492	0.00492	0.00492	0.00492	0.00492	0.00368	0.00368
53 wagn010616	0.05218	0.05366	0.05366	0.05366	0.05366	0.05366	0.05218	0.05218
54 wagn012969	0.05361	0.05509	0.05509	0.05509	0.05509	0.05509	0.05361	0.05361
55 wagn177351	0.05042	0.05189	0.05189	0.05189	0.05189	0.05189	0.05042	0.05042
56 podicUC243	0.10749	0.10930	0.10930	0.10930	0.10930	0.10930	0.10749	0.10749
57 podicUC244	0.10586	0.10767	0.10767	0.10767	0.10767	0.10767	0.10586	0.10586
58 podicUC278	0.10913	0.11095	0.11095	0.11095	0.11095	0.11095	0.10913	0.10913
59 podic53124	0.10849	0.11032	0.11032	0.11032	0.11032	0.11032	0.10849	0.10849
60 chaq186524	0.12481	0.12550	0.12550	0.12550	0.12550	0.12550	0.12481	0.12481
61 knuds13244	0.17003	0.17219	0.17219	0.17219	0.17219	0.17219	0.17003	0.17003



	17	18	19	20	21	22	23	24
17 Gren197007	-							
18 Gren197008	0.00000	-						
19 Gren197017	0.00122	0.00122	-					
20 Gren197044	0.00000	0.00000	0.00122	-				
21 StVn056421	0.00122	0.00122	0.00000	0.00122	-			
22 StVn056490	0.00122	0.00122	0.00000	0.00122	0.00000	-		
23 StVn056561	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000	-	
24 StVn056562	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000	0.00000	-
25 StVn056612	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000	0.00000	0.00000
26 StVn196895	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000	0.00000	0.00000
27 StVn196896	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
28 StVn196897	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
29 StVn196898	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000	0.00000	0.00000
30 StVn196899	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
31 StVn196900	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
32 Tobol86597	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000	0.00000	0.00000
33 Trin196726	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
34 Trin196727	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
35 Trin196728	0.00122	0.00122	0.00244	0.00122	0.00244	0.00244	0.00244	0.00244
36 Trin196729	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000	0.00000	0.00000
37 Trin196730	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
38 Trin196731	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
39 Trin196732	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
40 Trin196733	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
41 Trin196734	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
42 Trin196888	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
43 StVn056613	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000	0.00000	0.00000
44 StVn196894	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
45 Trin175424	0.00366	0.00366	0.00243	0.00366	0.00243	0.00243	0.00243	0.00243
46 Tobol86596	0.00122	0.00122	0.00000	0.00122	0.00000	0.00000	0.00000	0.00000
47 Trin196886	0.00244	0.00244	0.00122	0.00244	0.00122	0.00122	0.00122	0.00122
48 Trin175410	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
49 Trin175620	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
50 Trin196735	0.00000	0.00000	0.00122	0.00000	0.00122	0.00122	0.00122	0.00122
51 pall207682	0.00245	0.00245	0.00122	0.00245	0.00122	0.00122	0.00122	0.00122
52 pallWR8626	0.00492	0.00492	0.00368	0.00492	0.00368	0.00368	0.00368	0.00368
53 wagn010616	0.05366	0.05366	0.05218	0.05366	0.05218	0.05218	0.05218	0.05218
54 wagn012969	0.05509	0.05509	0.05361	0.05509	0.05361	0.05361	0.05361	0.05361
55 wagn177351	0.05189	0.05189	0.05042	0.05189	0.05042	0.05042	0.05042	0.05042
56 podicUC243	0.10930	0.10930	0.10749	0.10930	0.10749	0.10749	0.10749	0.10749
57 podicUC244	0.10767	0.10767	0.10586	0.10767	0.10586	0.10586	0.10586	0.10586
58 podicUC278	0.11095	0.11095	0.10913	0.11095	0.10913	0.10913	0.10913	0.10913
59 podic53124	0.11032	0.11032	0.10849	0.11032	0.10849	0.10849	0.10849	0.10849
60 chaq186524	0.12550	0.12550	0.12481	0.12550	0.12481	0.12481	0.12481	0.12481
61 knuds13244	0.17219	0.17219	0.17003	0.17219	0.17003	0.17003	0.17003	0.17003

	25	26	27	28	29	30	31	32
25 StVn056612	-							
26 StVn196895	0.00000	-						
27 StVn196896	0.00122	0.00122	-					
28 StVn196897	0.00122	0.00122	0.00000	-				
29 StVn196898	0.00000	0.00000	0.00122	0.00122	-			
30 StVn196899	0.00122	0.00122	0.00000	0.00000	0.00122	-		
31 StVn196900	0.00122	0.00122	0.00000	0.00000	0.00122	0.00000	-	
32 Tobo186597	0.00000	0.00000	0.00122	0.00122	0.00000	0.00122	0.00122	-
33 Trin196726	0.00122	0.00122	0.00000	0.00000	0.00122	0.00000	0.00000	0.00122
34 Trin196727	0.00122	0.00122	0.00000	0.00000	0.00122	0.00000	0.00000	0.00122
35 Trin196728	0.00244	0.00244	0.00122	0.00122	0.00244	0.00122	0.00122	0.00244
36 Trin196729	0.00000	0.00000	0.00122	0.00122	0.00000	0.00122	0.00122	0.00000
37 Trin196730	0.00122	0.00122	0.00000	0.00000	0.00122	0.00000	0.00000	0.00122
38 Trin196731	0.00122	0.00122	0.00000	0.00000	0.00122	0.00000	0.00000	0.00122
39 Trin196732	0.00122	0.00122	0.00000	0.00000	0.00122	0.00000	0.00000	0.00122
40 Trin196733	0.00122	0.00122	0.00000	0.00000	0.00122	0.00000	0.00000	0.00122
41 Trin196734	0.00122	0.00122	0.00000	0.00000	0.00122	0.00000	0.00000	0.00122
42 Trin196888	0.00122	0.00122	0.00000	0.00000	0.00122	0.00000	0.00000	0.00122
43 StVn056613	0.00000	0.00000	0.00122	0.00122	0.00000	0.00122	0.00122	0.00000
44 StVn196894	0.00122	0.00122	0.00000	0.00000	0.00122	0.00000	0.00000	0.00122
45 Trin175424	0.00243	0.00243	0.00366	0.00366	0.00243	0.00366	0.00366	0.00243
46 Tobo186596	0.00000	0.00000	0.00122	0.00122	0.00000	0.00122	0.00122	0.00000
47 Trin196886	0.00122	0.00122	0.00244	0.00244	0.00122	0.00244	0.00244	0.00122
48 Trin175410	0.00122	0.00122	0.00000	0.00000	0.00122	0.00000	0.00000	0.00122
49 Trin175620	0.00122	0.00122	0.00000	0.00000	0.00122	0.00000	0.00000	0.00122
50 Trin196735	0.00122	0.00122	0.00000	0.00000	0.00122	0.00000	0.00000	0.00122
51 pall207682	0.00122	0.00122	0.00245	0.00245	0.00122	0.00245	0.00245	0.00122
52 pallWR8626	0.00368	0.00368	0.00492	0.00492	0.00368	0.00492	0.00492	0.00368
53 wagn010616	0.05218	0.05218	0.05366	0.05366	0.05218	0.05366	0.05366	0.05218
54 wagn012969	0.05361	0.05361	0.05509	0.05509	0.05361	0.05509	0.05509	0.05361
55 wagn177351	0.05042	0.05042	0.05189	0.05189	0.05042	0.05189	0.05189	0.05042
56 podicUC243	0.10749	0.10749	0.10930	0.10930	0.10749	0.10930	0.10930	0.10749
57 podicUC244	0.10586	0.10586	0.10767	0.10767	0.10586	0.10767	0.10767	0.10586
58 podicUC278	0.10913	0.10913	0.11095	0.11095	0.10913	0.11095	0.11095	0.10913
59 podic53124	0.10849	0.10849	0.11032	0.11032	0.10849	0.11032	0.11032	0.10849
60 chaq186524	0.12481	0.12481	0.12550	0.12550	0.12481	0.12550	0.12550	0.12481
61 knuds13244	0.17003	0.17003	0.17219	0.17219	0.17003	0.17219	0.17219	0.17003

	33	34	35	36	37	38	39	40
33 Trin196726	-							
34 Trin196727	0.00000	-						
35 Trin196728	0.00122	0.00122	-					
36 Trin196729	0.00122	0.00122	0.00244	-				
37 Trin196730	0.00000	0.00000	0.00122	0.00122	-			
38 Trin196731	0.00000	0.00000	0.00122	0.00122	0.00000	-		
39 Trin196732	0.00000	0.00000	0.00122	0.00122	0.00000	0.00000	-	
40 Trin196733	0.00000	0.00000	0.00122	0.00122	0.00000	0.00000	0.00000	-
41 Trin196734	0.00000	0.00000	0.00122	0.00122	0.00000	0.00000	0.00000	0.00000
42 Trin196888	0.00000	0.00000	0.00122	0.00122	0.00000	0.00000	0.00000	0.00000
43 StVn056613	0.00122	0.00122	0.00244	0.00000	0.00122	0.00122	0.00122	0.00122
44 StVn196894	0.00000	0.00000	0.00122	0.00122	0.00000	0.00000	0.00000	0.00000
45 Trin175424	0.00366	0.00366	0.00490	0.00243	0.00366	0.00366	0.00366	0.00366
46 Tobol86596	0.00122	0.00122	0.00244	0.00000	0.00122	0.00122	0.00122	0.00122
47 Trin196886	0.00244	0.00244	0.00368	0.00122	0.00244	0.00244	0.00244	0.00244
48 Trin175410	0.00000	0.00000	0.00122	0.00122	0.00000	0.00000	0.00000	0.00000
49 Trin175620	0.00000	0.00000	0.00122	0.00122	0.00000	0.00000	0.00000	0.00000
50 Trin196735	0.00000	0.00000	0.00122	0.00122	0.00000	0.00000	0.00000	0.00000
51 pall207682	0.00245	0.00245	0.00367	0.00122	0.00245	0.00245	0.00245	0.00245
52 pallWR8626	0.00492	0.00492	0.00618	0.00368	0.00492	0.00492	0.00492	0.00492
53 wagn010616	0.05366	0.05366	0.05546	0.05218	0.05366	0.05366	0.05366	0.05366
54 wagn012969	0.05509	0.05509	0.05691	0.05361	0.05509	0.05509	0.05509	0.05509
55 wagn177351	0.05189	0.05189	0.05367	0.05042	0.05189	0.05189	0.05189	0.05189
56 podicUC243	0.10930	0.10930	0.10910	0.10749	0.10930	0.10930	0.10930	0.10930
57 podicUC244	0.10767	0.10767	0.10747	0.10586	0.10767	0.10767	0.10767	0.10767
58 podicUC278	0.11095	0.11095	0.11075	0.10913	0.11095	0.11095	0.11095	0.11095
59 podic53124	0.11032	0.11032	0.11010	0.10849	0.11032	0.11032	0.11032	0.11032
60 chaq186524	0.12550	0.12550	0.12809	0.12481	0.12550	0.12550	0.12550	0.12550
61 knuds13244	0.17219	0.17219	0.16926	0.17003	0.17219	0.17219	0.17219	0.17219

	41	42	43	44	45	46	47	48
41 Trin196734	-							
42 Trin196888	0.00000	-						
43 StVn056613	0.00122	0.00122	-					
44 StVn196894	0.00000	0.00000	0.00122	-				
45 Trin175424	0.00366	0.00366	0.00243	0.00366	-			
46 Tobo186596	0.00122	0.00122	0.00000	0.00122	0.00243	-		
47 Trin196886	0.00244	0.00244	0.00122	0.00244	0.00367	0.00122	-	
48 Trin175410	0.00000	0.00000	0.00122	0.00000	0.00366	0.00122	0.00244	-
49 Trin175620	0.00000	0.00000	0.00122	0.00000	0.00366	0.00122	0.00244	0.00000
50 Trin196735	0.00000	0.00000	0.00122	0.00000	0.00366	0.00122	0.00244	0.00000
51 pall207682	0.00245	0.00245	0.00122	0.00245	0.00366	0.00122	0.00244	0.00245
52 pallWR8626	0.00492	0.00492	0.00368	0.00492	0.00616	0.00368	0.00494	0.00492
53 wagn010616	0.05366	0.05366	0.05218	0.05366	0.05537	0.05218	0.05399	0.05366
54 wagn012969	0.05509	0.05509	0.05361	0.05509	0.05681	0.05361	0.05542	0.05509
55 wagn177351	0.05189	0.05189	0.05042	0.05189	0.05358	0.05042	0.05220	0.05189
56 podicUC243	0.10930	0.10930	0.10749	0.10930	0.10679	0.10749	0.10987	0.10930
57 podicUC244	0.10767	0.10767	0.10586	0.10767	0.10514	0.10586	0.10825	0.10767
58 podicUC278	0.11095	0.11095	0.10913	0.11095	0.10845	0.10913	0.11152	0.11095
59 podic53124	0.11032	0.11032	0.10849	0.11032	0.10769	0.10849	0.11093	0.11032
60 chaq186524	0.12550	0.12550	0.12481	0.12550	0.12406	0.12481	0.12231	0.12550
61 knuds13244	0.17219	0.17219	0.17003	0.17219	0.17504	0.17003	0.17300	0.17219

	49	50	51	52	53	54	55	56
49 Trin175620	-							
50 Trin196735	0.00000	-						
51 pall207682	0.00245	0.00245	-					
52 pallWR8626	0.00492	0.00492	0.00245	-				
53 wagn010616	0.05366	0.05366	0.05211	0.05210	-			
54 wagn012969	0.05509	0.05509	0.05353	0.05716	0.01520	-		
55 wagn177351	0.05189	0.05189	0.05036	0.05035	0.00244	0.01512	-	
56 podicUC243	0.10930	0.10930	0.10678	0.10683	0.10738	0.11043	0.10985	-
57 podicUC244	0.10767	0.10767	0.10515	0.10520	0.10577	0.10879	0.10824	0.00122
58 podicUC278	0.11095	0.11095	0.10843	0.10848	0.10577	0.11210	0.10824	0.00629
59 podic53124	0.11032	0.11032	0.10773	0.10778	0.10532	0.11143	0.10787	0.00746
60 chaq186524	0.12550	0.12550	0.12680	0.12192	0.10948	0.11809	0.11008	0.12356
61 knuds13244	0.17219	0.17219	0.16998	0.17006	0.15841	0.16599	0.16154	0.15559

	57	58	59	60	61
57 podicUC244	-				
58 podicUC278	0.00499	-			
59 podic53124	0.00618	0.00875	-		
60 chaq186524	0.12184	0.12715	0.12299	-	
61 knuds13244	0.15354	0.15570	0.15109	0.12095	-

Table V. Absolute base differences among samples for the 12S dataset.

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
1 Gren006881	-														
2 Gren006882	0	-													
3 Gren006883	0	0	-												
4 Gren006939	0	0	0	-											
5 Gren196977	1	1	1	1	-										
6 Gren196978	0	0	0	0	1	-									
7 Gren196979	1	1	1	1	0	1	-								
8 Gren196980	1	1	1	1	0	1	0	-							
9 Gren196999	0	0	0	0	1	0	1	1	-						
10 Gren197000	1	1	1	1	0	1	0	0	1	-					
11 Gren197001	1	1	1	1	0	1	0	0	1	0	-				
12 Gren197002	1	1	1	1	0	1	0	0	1	0	0	-			
13 Gren197003	1	1	1	1	0	1	0	0	1	0	0	0	-		
14 Gren197004	1	1	1	1	0	1	0	0	1	0	0	0	0	-	
15 Gren197005	0	0	0	0	1	0	1	1	0	1	1	1	1	1	-
16 Gren197006	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
17 Gren197007	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
18 Gren197008	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
19 Gren197017	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
20 Gren197044	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
21 StVn056421	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
22 StVn056490	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
23 StVn056561	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
24 StVn056562	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
25 StVn056612	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
26 StVn196895	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
27 StVn196896	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
28 StVn196897	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
29 StVn196898	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
30 StVn196899	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
31 StVn196900	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
32 Tobol86597	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
33 Trinl96726	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
34 Trinl96727	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
35 Trinl96728	2	2	2	2	1	2	1	1	2	1	1	1	1	1	2
36 Trinl96729	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
37 Trinl96730	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
38 Trinl96731	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
39 Trinl96732	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
40 Trinl96733	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
41 Trinl96734	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
42 Trinl96888	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
43 StVn056613	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
44 StVn196894	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
45 Trinl75424	2	2	2	2	3	2	3	3	2	3	3	3	3	3	2
46 Tobol86596	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
47 Trinl96886	1	1	1	1	2	1	2	2	1	2	2	2	2	2	1
48 Trinl75410	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
49 Trinl75620	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
50 Trinl96735	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
51 pall1207682	1	1	1	1	2	1	2	2	1	2	2	2	2	2	1
52 pallWR8626	3	3	3	3	4	3	4	4	3	4	4	4	4	4	3
53 wagn010616	37	37	37	37	38	37	38	38	37	38	38	38	38	38	37
54 wagn012969	38	38	38	38	39	38	39	39	38	39	39	39	39	39	38
55 wagn177351	36	36	36	36	37	36	37	37	36	37	37	37	37	37	36
56 podicUC243	68	68	68	68	69	68	69	69	68	69	69	69	69	69	68
57 podicUC244	67	67	67	67	68	67	68	68	67	68	68	68	68	68	67
58 podicUC278	69	69	69	69	70	69	70	70	69	70	70	70	70	70	69
59 podic53124	68	68	68	68	69	68	69	69	68	69	69	69	69	69	68
60 chaq186524	78	78	78	78	78	78	78	78	78	78	78	78	78	78	78
61 knuds13244	99	99	99	99	100	99	100	100	99	100	100	100	100	100	99

	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30
16 Gren197006	-														
17 Gren197007	1	-													
18 Gren197008	1	0	-												
19 Gren197017	0	1	1	-											
20 Gren197044	1	0	0	1	-										
21 StVn056421	0	1	1	0	1	-									
22 StVn056490	0	1	1	0	1	0	-								
23 StVn056561	0	1	1	0	1	0	0	-							
24 StVn056562	0	1	1	0	1	0	0	0	-						
25 StVn056612	0	1	1	0	1	0	0	0	0	-					
26 StVn196895	0	1	1	0	1	0	0	0	0	0	-				
27 StVn196896	1	0	0	1	0	1	1	1	1	1	1	-			
28 StVn196897	1	0	0	1	0	1	1	1	1	1	1	0	-		
29 StVn196898	0	1	1	0	1	0	0	0	0	0	0	1	1	-	
30 StVn196899	1	0	0	1	0	1	1	1	1	1	1	0	0	1	-
31 StVn196900	1	0	0	1	0	1	1	1	1	1	1	0	0	1	0
32 Tobol86597	0	1	1	0	1	0	0	0	0	0	0	1	1	0	1
33 Trin196726	1	0	0	1	0	1	1	1	1	1	1	0	0	1	0
34 Trin196727	1	0	0	1	0	1	1	1	1	1	1	0	0	1	0
35 Trin196728	2	1	1	2	1	2	2	2	2	2	2	1	1	2	1
36 Trin196729	0	1	1	0	1	0	0	0	0	0	0	1	1	0	1
37 Trin196730	1	0	0	1	0	1	1	1	1	1	1	0	0	1	0
38 Trin196731	1	0	0	1	0	1	1	1	1	1	1	0	0	1	0
39 Trin196732	1	0	0	1	0	1	1	1	1	1	1	0	0	1	0
40 Trin196733	1	0	0	1	0	1	1	1	1	1	1	0	0	1	0
41 Trin196734	1	0	0	1	0	1	1	1	1	1	1	0	0	1	0
42 Trin196888	1	0	0	1	0	1	1	1	1	1	1	0	0	1	0
43 StVn056613	0	1	1	0	1	0	0	0	0	0	0	1	1	0	1
44 StVn196894	1	0	0	1	0	1	1	1	1	1	1	0	0	1	0
45 Trin175424	2	3	3	2	3	2	2	2	2	2	2	3	3	2	3
46 Tobol86596	0	1	1	0	1	0	0	0	0	0	0	1	1	0	1
47 Trin196886	1	2	2	1	2	1	1	1	1	1	1	2	2	1	2
48 Trin175410	1	0	0	1	0	1	1	1	1	1	1	0	0	1	0
49 Trin175620	1	0	0	1	0	1	1	1	1	1	1	0	0	1	0
50 Trin196735	1	0	0	1	0	1	1	1	1	1	1	0	0	1	0
51 pall207682	1	2	2	1	2	1	1	1	1	1	1	2	2	1	2
52 pallWR8626	3	4	4	3	4	3	3	3	3	3	3	4	4	3	4
53 wagn010616	37	38	38	37	38	37	37	37	37	37	37	38	38	37	38
54 wagn012969	38	39	39	38	39	38	38	38	38	38	38	39	39	38	39
55 wagn177351	36	37	37	36	37	36	36	36	36	36	36	37	37	36	37
56 podicUC243	68	69	69	68	69	68	68	68	68	68	68	69	69	68	69
57 podicUC244	67	68	68	67	68	67	67	67	67	67	67	68	68	67	68
58 podicUC278	69	70	70	69	70	69	69	69	69	69	69	70	70	69	70
59 podic53124	68	69	69	68	69	68	68	68	68	68	68	69	69	68	69
60 chaq186524	78	78	78	78	78	78	78	78	78	78	78	78	78	78	78
61 knuds13244	99	100	100	99	100	99	99	99	99	99	99	100	100	99	100

	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45
31 StVn196900	-														
32 Tobo186597	1	-													
33 Trin196726	0	1	-												
34 Trin196727	0	1	0	-											
35 Trin196728	1	2	1	1	-										
36 Trin196729	1	0	1	1	2	-									
37 Trin196730	0	1	0	0	1	1	-								
38 Trin196731	0	1	0	0	1	1	0	-							
39 Trin196732	0	1	0	0	1	1	0	0	-						
40 Trin196733	0	1	0	0	1	1	0	0	0	-					
41 Trin196734	0	1	0	0	1	1	0	0	0	0	-				
42 Trin196888	0	1	0	0	1	1	0	0	0	0	0	-			
43 StVn056613	1	0	1	1	2	0	1	1	1	1	1	1	-		
44 StVn196894	0	1	0	0	1	1	0	0	0	0	0	0	1	-	
45 Trin175424	3	2	3	3	4	2	3	3	3	3	3	3	2	3	-
46 Tobo186596	1	0	1	1	2	0	1	1	1	1	1	1	0	1	2
47 Trin196886	2	1	2	2	3	1	2	2	2	2	2	2	1	2	3
48 Trin175410	0	1	0	0	1	1	0	0	0	0	0	0	1	0	3
49 Trin175620	0	1	0	0	1	1	0	0	0	0	0	0	1	0	3
50 Trin196735	0	1	0	0	1	1	0	0	0	0	0	0	1	0	3
51 pall207682	2	1	2	2	3	1	2	2	2	2	2	2	1	2	3
52 pallWR8626	4	3	4	4	5	3	4	4	4	4	4	4	3	4	5
53 wagn010616	38	37	38	38	39	37	38	38	38	38	38	38	37	38	39
54 wagn012969	39	38	39	39	40	38	39	39	39	39	39	39	38	39	40
55 wagn177351	37	36	37	37	38	36	37	37	37	37	37	37	36	37	38
56 podicUC243	69	68	69	69	69	68	69	69	69	69	69	69	68	69	68
57 podicUC244	68	67	68	68	68	67	68	68	68	68	68	68	67	68	67
58 podicUC278	70	69	70	70	70	69	70	70	70	70	70	70	69	70	69
59 podic53124	69	68	69	69	69	68	69	69	69	69	69	69	68	69	68
60 chaq186524	78	78	78	78	79	78	78	78	78	78	78	78	78	78	78
61 knuds13244	100	99	100	100	99	99	100	100	100	100	100	100	99	100	101

	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60
46 Tobo186596	-														
47 Trin196886	1	-													
48 Trin175410	1	2	-												
49 Trin175620	1	2	0	-											
50 Trin196735	1	2	0	0	-										
51 pall207682	1	2	2	2	2	-									
52 pallWR8626	3	4	4	4	4	2	-								
53 wagn010616	37	38	38	38	38	37	37	-							
54 wagn012969	38	39	39	39	39	38	40	12	-						
55 wagn177351	36	37	37	37	37	36	36	2	12	-					
56 podicUC243	68	69	69	69	69	68	68	68	70	69	-				
57 podicUC244	67	68	68	68	68	67	67	67	69	68	1	-			
58 podicUC278	69	70	70	70	70	69	69	67	71	68	5	4	-		
59 podic53124	68	69	69	69	69	68	68	66	70	67	6	5	7	-	
60 chaq186524	78	77	78	78	78	79	77	71	75	71	77	76	79	77	-
61 knuds13244	99	100	100	100	100	99	99	94	98	95	95	94	95	93	78

61 knuds13244 -

Table VI. 16S rDNA sequence haplotypes for *L. validus* samples from: Grenada (Gren), St. Vincent (StVn), Tobago (Tobo), and Trinidad (Trin). Samples A—C represent multiple samples. Asterisks indicate samples used to represent a haplotype. Haplotypes D—G each consist of a single sample: (D) Trin196886, (E) Trin175410, (F) Trin175620, and (G) Trin196735.

A	B	C
Gren006881	Trin175424	*Tobo186596
Gren006882	*Trin196726	Tobo186597
Gren006883	Trin196727	
Gren006939	Trin196728	
Gren196977	Trin196729	
Gren196978	Trin196730	
Gren196979	Trin196731	
Gren196980	Trin196732	
Gren196999	Trin196733	
Gren197000	Trin196734	
Gren197001	Trin196888	
Gren197002		
Gren197003		
Gren197004		
Gren197005		
Gren197006		
Gren197007		
Gren197008		
Gren197017		
Gren197044		
StVn056421		
StVn056490		
StVn056561		
StVn056562		
StVn056612		
*StVn056613		
StVn196894		
StVn196895		
StVn196896		
StVn196897		
StVn196898		
StVn196899		
StVn196900		



Table VII. Base composition percentages of 16S mtDNA gene sequences (averages among samples from each taxa).

	<i>validus</i>	<i>pallidirostris</i>	<i>podicipinus</i>	<i>wagneri</i>	<i>chaquensis</i>	<i>knudseni</i>	All taxa
A	0.34790	0.34899	0.34436	0.35052	0.35491	0.34591	0.34791
T	0.23839	0.23906	0.23703	0.23859	0.25747	0.23772	0.23863
G	0.17936	0.17859	0.17840	0.17752	0.18065	0.17438	0.17913
C	0.23435	0.23337	0.24024	0.23336	0.20697	0.24199	0.23433









	33	34	35	36	37	38	39	40
33 Trin196726	-							
34 Trin196727	0.00000	-						
35 Trin196728	0.00000	0.00000	-					
36 Trin196729	0.00000	0.00000	0.00000	-				
37 Trin196730	0.00000	0.00000	0.00000	0.00000	-			
38 Trin196731	0.00000	0.00000	0.00000	0.00000	0.00000	-		
39 Trin196732	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000	-	
40 Trin196733	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000	-
41 Trin196734	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000
42 Trin196888	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000
43 StVn056613	0.00215	0.00215	0.00215	0.00215	0.00215	0.00215	0.00215	0.00215
44 StVn196894	0.00215	0.00215	0.00215	0.00215	0.00215	0.00215	0.00215	0.00215
45 Trin175424	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000
46 Tobo186596	0.00143	0.00143	0.00143	0.00143	0.00143	0.00143	0.00143	0.00143
47 Trin196886	0.00144	0.00144	0.00144	0.00144	0.00144	0.00144	0.00144	0.00144
48 Trin175410	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000
49 Trin175620	0.00071	0.00071	0.00071	0.00071	0.00071	0.00071	0.00071	0.00071
50 Trin196735	0.00071	0.00071	0.00071	0.00071	0.00071	0.00071	0.00071	0.00071
51 pall1207682	0.00143	0.00143	0.00143	0.00143	0.00143	0.00143	0.00143	0.00143
52 pallWR8626	0.00506	0.00506	0.00506	0.00506	0.00506	0.00506	0.00506	0.00506
53 wagn010616	0.09245	0.09245	0.09245	0.09245	0.09245	0.09245	0.09245	0.09245
54 wagn012969	0.09373	0.09373	0.09373	0.09373	0.09373	0.09373	0.09373	0.09373
55 wagn177351	0.09569	0.09569	0.09569	0.09569	0.09569	0.09569	0.09569	0.09569
56 podicUC243	0.15544	0.15544	0.15544	0.15544	0.15544	0.15544	0.15544	0.15544
57 podicUC244	0.15544	0.15544	0.15544	0.15544	0.15544	0.15544	0.15544	0.15544
58 podicUC278	0.16514	0.16514	0.16514	0.16514	0.16514	0.16514	0.16514	0.16514
59 podic53124	0.15941	0.15941	0.15941	0.15941	0.15941	0.15941	0.15941	0.15941
60 chaq186524	0.20757	0.20757	0.20757	0.20757	0.20757	0.20757	0.20757	0.20757
61 knuds13244	0.26297	0.26297	0.26297	0.26297	0.26297	0.26297	0.26297	0.26297

	41	42	43	44	45	46	47	48
41 Trin196734	-							
42 Trin196888	0.00000	-						
43 StVn056613	0.00215	0.00215	-					
44 StVn196894	0.00215	0.00215	0.00000	-				
45 Trin175424	0.00000	0.00000	0.00215	0.00215	-			
46 Tobo186596	0.00143	0.00143	0.00359	0.00359	0.00143	-		
47 Trin196886	0.00144	0.00144	0.00359	0.00359	0.00144	0.00288	-	
48 Trin175410	0.00000	0.00000	0.00215	0.00215	0.00000	0.00143	0.00144	-
49 Trin175620	0.00071	0.00071	0.00287	0.00287	0.00071	0.00215	0.00216	0.00071
50 Trin196735	0.00071	0.00071	0.00287	0.00287	0.00071	0.00215	0.00215	0.00072
51 pall1207682	0.00143	0.00143	0.00360	0.00360	0.00143	0.00287	0.00287	0.00143
52 pallWR8626	0.00506	0.00506	0.00725	0.00725	0.00506	0.00653	0.00655	0.00506
53 wagn010616	0.09245	0.09245	0.09590	0.09590	0.09245	0.09472	0.09245	0.09248
54 wagn012969	0.09373	0.09373	0.09723	0.09723	0.09373	0.09603	0.09373	0.09376
55 wagn177351	0.09569	0.09569	0.09918	0.09918	0.09569	0.09799	0.09569	0.09572
56 podicUC243	0.15544	0.15544	0.15815	0.15815	0.15544	0.15602	0.15222	0.15445
57 podicUC244	0.15544	0.15544	0.15815	0.15815	0.15544	0.15602	0.15222	0.15445
58 podicUC278	0.16514	0.16514	0.16794	0.16794	0.16514	0.16580	0.16172	0.16413
59 podic53124	0.15941	0.15941	0.16222	0.16222	0.15941	0.16000	0.15612	0.15842
60 chaq186524	0.20757	0.20757	0.20742	0.20742	0.20757	0.20429	0.20961	0.20771
61 knuds13244	0.26297	0.26297	0.26754	0.26754	0.26297	0.26373	0.26098	0.26319

	49	50	51	52	53	54	55	56
49 Trin175620	-							
50 Trin196735	0.00143	-						
51 pall207682	0.00215	0.00215	-					
52 pallWR8626	0.00580	0.00578	0.00506	-				
53 wagn010616	0.09116	0.09344	0.09252	0.08836	-			
54 wagn012969	0.09513	0.09471	0.09380	0.08959	0.01731	-		
55 wagn177351	0.09438	0.09668	0.09574	0.09153	0.00215	0.01655	-	
56 podicUC243	0.15283	0.15672	0.15462	0.15340	0.14229	0.13983	0.14388	-
57 podicUC244	0.15283	0.15672	0.15462	0.15340	0.14229	0.13983	0.14388	0.00000
58 podicUC278	0.16241	0.16470	0.16430	0.16301	0.14912	0.14805	0.15081	0.00879
59 podic53124	0.15677	0.15897	0.15859	0.15736	0.15219	0.14994	0.15386	0.01954
60 chaq186524	0.20559	0.20620	0.20792	0.20196	0.21247	0.22312	0.21427	0.17661
61 knuds13244	0.26102	0.26460	0.26618	0.25933	0.28386	0.28525	0.28761	0.26021

	57	58	59	60	61
57 podicUC244	-				
58 podicUC278	0.00879	-			
59 podic53124	0.01954	0.02104	-		
60 chaq186524	0.17661	0.17904	0.17989	-	
61 knuds13244	0.26021	0.26906	0.26391	0.20697	-







	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45
31 StVn196900	-														
32 Tobol86597	5	-													
33 Trin196726	3	2	-												
34 Trin196727	3	2	0	-											
35 Trin196728	3	2	0	0	-										
36 Trin196729	3	2	0	0	0	-									
37 Trin196730	3	2	0	0	0	0	-								
38 Trin196731	3	2	0	0	0	0	0	-							
39 Trin196732	3	2	0	0	0	0	0	0	-						
40 Trin196733	3	2	0	0	0	0	0	0	0	-					
41 Trin196734	3	2	0	0	0	0	0	0	0	0	-				
42 Trin196888	3	2	0	0	0	0	0	0	0	0	0	-			
43 StVn056613	0	5	3	3	3	3	3	3	3	3	3	3	-		
44 StVn196894	0	5	3	3	3	3	3	3	3	3	3	3	0	-	
45 Trin175424	3	2	0	0	0	0	0	0	0	0	0	0	3	3	-
46 Tobol86596	5	0	2	2	2	2	2	2	2	2	2	2	5	5	2
47 Trin196886	5	4	2	2	2	2	2	2	2	2	2	2	5	5	2
48 Trin175410	3	2	0	0	0	0	0	0	0	0	0	0	3	3	0
49 Trin175620	4	3	1	1	1	1	1	1	1	1	1	1	4	4	1
50 Trin196735	4	3	1	1	1	1	1	1	1	1	1	1	4	4	1
51 pall207682	5	4	2	2	2	2	2	2	2	2	2	2	5	5	2
52 pallWR8626	10	9	7	7	7	7	7	7	7	7	7	7	10	10	7
53 wagn010616	105	104	102	102	102	102	102	102	102	102	102	102	105	105	102
54 wagn012969	105	104	102	102	102	102	102	102	102	102	102	102	105	105	102
55 wagn177351	108	107	105	105	105	105	105	105	105	105	105	105	108	108	105
56 podicUC243	160	158	158	158	158	158	158	158	158	158	158	158	160	160	158
57 podicUC244	160	158	158	158	158	158	158	158	158	158	158	158	160	160	158
58 podicUC278	166	164	164	164	164	164	164	164	164	164	164	164	166	166	164
59 podic53124	163	161	161	161	161	161	161	161	161	161	161	161	163	163	161
60 chaq186524	191	189	191	191	191	191	191	191	191	191	191	191	191	191	191
61 knuds13244	236	233	233	233	233	233	233	233	233	233	233	233	236	236	233

	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60
46 Tobol86596	-														
47 Trin196886	4	-													
48 Trin175410	2	2	-												
49 Trin175620	3	3	1	-											
50 Trin196735	3	3	1	2	-										
51 pall207682	4	4	2	3	3	-									
52 pallWR8626	9	9	7	8	8	7	-								
53 wagn010616	104	102	102	101	103	102	98	-							
54 wagn012969	104	102	102	103	103	102	98	23	-						
55 wagn177351	107	105	105	104	106	105	101	3	22	-					
56 podicUC243	158	156	157	156	159	157	156	147	143	148	-				
57 podicUC244	158	156	157	156	159	157	156	147	143	148	0	-			
58 podicUC278	164	162	163	162	164	163	162	151	148	152	12	12	-		
59 podic53124	161	159	160	159	161	160	159	154	150	155	26	26	28	-	
60 chaq186524	189	192	191	190	190	191	187	192	198	193	172	172	174	175	-
61 knuds13244	233	232	233	232	234	235	231	242	242	244	230	230	235	233	196

61

61 knuds13244 -

Table X. Combined sequence haplotypes for *L. validus* samples from: Grenada (Gren), St. Vincent (StVn), Tobago (Tobo), and Trinidad (Trin). Haplotypes A–D represent multiple samples. Samples used to represent each haplotype are indicated by an asterisk. Haplotypes E–K each consist of a single sample: (E) Trin175410, (F) Trin175424, (G) Trin175620, (H) Trin196728, (I) Trin196729, (J) Trin196735, and (K) Trin196886.

A	B	C	D
Gren006881	Gren196977	*Trin196726	*Tobo186596
Gren006882	Gren196979	Trin196727	Tobo186597
Gren006883	Gren196980	Trin196730	
Gren006939	Gren197000	Trin196731	
Gren196978	Gren197001	Trin196732	
Gren196999	Gren197002	Trin196733	
Gren197005	Gren197003	Trin196734	
Gren197006	Gren197004	Trin196888	
Gren197017	Gren197007		
StVn056421	Gren197008		
StVn056490	Gren197044		
StVn056561	StVn196894		
StVn056562	*StVn196896		
StVn056612	StVn196897		
*StVn056613	StVn196899		
StVn196895	StVn196900		
StVn196898			

Table XI. Pair-wise genetic distances for the combined dataset. Distances corrected using the general time-reversible model with gamma distributed rates for variable sites.

	1	2	3	4	5	6	7	8
1 Gren006881	-							
2 Gren006882	0.00000	-						
3 Gren006883	0.00000	0.00000	-					
4 Gren006939	0.00000	0.00000	0.00000	-				
5 Gren196977	0.00045	0.00045	0.00045	0.00045	-			
6 Gren196978	0.00000	0.00000	0.00000	0.00000	0.00045	-		
7 Gren196979	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	-	
8 Gren196980	0.00045	0.00045	0.00045	0.00045	0.00045	0.00000	0.00000	-
9 Gren196999	0.00000	0.00000	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045
10 Gren197000	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000
11 Gren197001	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000
12 Gren197002	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000
13 Gren197003	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000
14 Gren197004	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000
15 Gren197005	0.00000	0.00000	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045
16 Gren197006	0.00000	0.00000	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045
17 Gren197007	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000
18 Gren197008	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000
19 Gren197017	0.00000	0.00000	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045
20 Gren197044	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000
21 StVn056421	0.00000	0.00000	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045
22 StVn056490	0.00000	0.00000	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045
23 StVn056561	0.00000	0.00000	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045
24 StVn056562	0.00000	0.00000	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045
25 StVn056612	0.00000	0.00000	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045
26 StVn196895	0.00000	0.00000	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045
27 StVn196896	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000
28 StVn196897	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000
29 StVn196898	0.00000	0.00000	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045
30 StVn196899	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000
31 StVn196900	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000
32 Tobo186597	0.00225	0.00225	0.00225	0.00225	0.00271	0.00225	0.00271	0.00271
33 Trin196726	0.00180	0.00180	0.00180	0.00180	0.00135	0.00180	0.00135	0.00135
34 Trin196727	0.00180	0.00180	0.00180	0.00180	0.00135	0.00180	0.00135	0.00135
35 Trin196728	0.00225	0.00225	0.00225	0.00225	0.00180	0.00225	0.00180	0.00180
36 Trin196729	0.00135	0.00135	0.00135	0.00135	0.00180	0.00135	0.00180	0.00180
37 Trin196730	0.00180	0.00180	0.00180	0.00180	0.00135	0.00180	0.00135	0.00135
38 Trin196731	0.00180	0.00180	0.00180	0.00180	0.00135	0.00180	0.00135	0.00135
39 Trin196732	0.00180	0.00180	0.00180	0.00180	0.00135	0.00180	0.00135	0.00135
40 Trin196733	0.00180	0.00180	0.00180	0.00180	0.00135	0.00180	0.00135	0.00135
41 Trin196734	0.00180	0.00180	0.00180	0.00180	0.00135	0.00180	0.00135	0.00135
42 Trin196888	0.00180	0.00180	0.00180	0.00180	0.00135	0.00180	0.00135	0.00135
43 StVn056613	0.00000	0.00000	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045
44 StVn196894	0.00045	0.00045	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000
45 Trin175424	0.00225	0.00225	0.00225	0.00225	0.00271	0.00225	0.00271	0.00271
46 Tobo186596	0.00225	0.00225	0.00225	0.00225	0.00271	0.00225	0.00271	0.00271
47 Trin196886	0.00271	0.00271	0.00271	0.00271	0.00316	0.00271	0.00316	0.00316
48 Trin175410	0.00180	0.00180	0.00180	0.00180	0.00135	0.00180	0.00135	0.00135
49 Trin175620	0.00225	0.00225	0.00225	0.00225	0.00180	0.00225	0.00180	0.00180
50 Trin196735	0.00226	0.00226	0.00226	0.00226	0.00180	0.00226	0.00180	0.00180
51 pall207682	0.00271	0.00271	0.00271	0.00271	0.00317	0.00271	0.00317	0.00317
52 pallWR8626	0.00591	0.00591	0.00591	0.00591	0.00637	0.00591	0.00637	0.00637
53 wagn010616	0.07859	0.07859	0.07859	0.07859	0.07917	0.07859	0.07917	0.07917
54 wagn012969	0.07983	0.07983	0.07983	0.07983	0.08041	0.07983	0.08041	0.08041
55 wagn177351	0.07974	0.07974	0.07974	0.07974	0.08032	0.07974	0.08032	0.08032
56 podicUC243	0.13826	0.13826	0.13826	0.13826	0.13899	0.13826	0.13899	0.13899
57 podicUC244	0.13761	0.13761	0.13761	0.13761	0.13834	0.13761	0.13834	0.13834
58 podicUC278	0.14460	0.14460	0.14460	0.14460	0.14535	0.14460	0.14535	0.14535
59 podic53124	0.14119	0.14119	0.14119	0.14119	0.14194	0.14119	0.14194	0.14194
60 chaq186524	0.17386	0.17386	0.17386	0.17386	0.17424	0.17386	0.17424	0.17424
61 knuds13244	0.22738	0.22738	0.22738	0.22738	0.22830	0.22738	0.22830	0.22830

	9	10	11	12	13	14	15	16
9 Gren196999	-							
10 Gren197000	0.00045	-						
11 Gren197001	0.00045	0.00000	-					
12 Gren197002	0.00045	0.00000	0.00000	-				
13 Gren197003	0.00045	0.00000	0.00000	0.00000	-			
14 Gren197004	0.00045	0.00000	0.00000	0.00000	0.00000	-		
15 Gren197005	0.00000	0.00045	0.00045	0.00045	0.00045	0.00045	-	
16 Gren197006	0.00000	0.00045	0.00045	0.00045	0.00045	0.00045	0.00000	-
17 Gren197007	0.00045	0.00000	0.00000	0.00000	0.00000	0.00000	0.00045	0.00045
18 Gren197008	0.00045	0.00000	0.00000	0.00000	0.00000	0.00000	0.00045	0.00045
19 Gren197017	0.00000	0.00045	0.00045	0.00045	0.00045	0.00045	0.00000	0.00000
20 Gren197044	0.00045	0.00000	0.00000	0.00000	0.00000	0.00000	0.00045	0.00045
21 StVn056421	0.00000	0.00045	0.00045	0.00045	0.00045	0.00045	0.00000	0.00000
22 StVn056490	0.00000	0.00045	0.00045	0.00045	0.00045	0.00045	0.00000	0.00000
23 StVn056561	0.00000	0.00045	0.00045	0.00045	0.00045	0.00045	0.00000	0.00000
24 StVn056562	0.00000	0.00045	0.00045	0.00045	0.00045	0.00045	0.00000	0.00000
25 StVn056612	0.00000	0.00045	0.00045	0.00045	0.00045	0.00045	0.00000	0.00000
26 StVn196895	0.00000	0.00045	0.00045	0.00045	0.00045	0.00045	0.00000	0.00000
27 StVn196896	0.00045	0.00000	0.00000	0.00000	0.00000	0.00000	0.00045	0.00045
28 StVn196897	0.00045	0.00000	0.00000	0.00000	0.00000	0.00000	0.00045	0.00045
29 StVn196898	0.00000	0.00045	0.00045	0.00045	0.00045	0.00045	0.00000	0.00000
30 StVn196899	0.00045	0.00000	0.00000	0.00000	0.00000	0.00000	0.00045	0.00045
31 StVn196900	0.00045	0.00000	0.00000	0.00000	0.00000	0.00000	0.00045	0.00045
32 Tobo186597	0.00225	0.00271	0.00271	0.00271	0.00271	0.00271	0.00225	0.00225
33 Trin196726	0.00180	0.00135	0.00135	0.00135	0.00135	0.00135	0.00180	0.00180
34 Trin196727	0.00180	0.00135	0.00135	0.00135	0.00135	0.00135	0.00180	0.00180
35 Trin196728	0.00225	0.00180	0.00180	0.00180	0.00180	0.00180	0.00225	0.00225
36 Trin196729	0.00135	0.00180	0.00180	0.00180	0.00180	0.00180	0.00135	0.00135
37 Trin196730	0.00180	0.00135	0.00135	0.00135	0.00135	0.00135	0.00180	0.00180
38 Trin196731	0.00180	0.00135	0.00135	0.00135	0.00135	0.00135	0.00180	0.00180
39 Trin196732	0.00180	0.00135	0.00135	0.00135	0.00135	0.00135	0.00180	0.00180
40 Trin196733	0.00180	0.00135	0.00135	0.00135	0.00135	0.00135	0.00180	0.00180
41 Trin196734	0.00180	0.00135	0.00135	0.00135	0.00135	0.00135	0.00180	0.00180
42 Trin196888	0.00180	0.00135	0.00135	0.00135	0.00135	0.00135	0.00180	0.00180
43 StVn056613	0.00000	0.00045	0.00045	0.00045	0.00045	0.00045	0.00000	0.00000
44 StVn196894	0.00045	0.00000	0.00000	0.00000	0.00000	0.00000	0.00045	0.00045
45 Trin175424	0.00225	0.00271	0.00271	0.00271	0.00271	0.00271	0.00225	0.00225
46 Tobo186596	0.00225	0.00271	0.00271	0.00271	0.00271	0.00271	0.00225	0.00225
47 Trin196886	0.00271	0.00316	0.00316	0.00316	0.00316	0.00316	0.00271	0.00271
48 Trin175410	0.00180	0.00135	0.00135	0.00135	0.00135	0.00135	0.00180	0.00180
49 Trin175620	0.00225	0.00180	0.00180	0.00180	0.00180	0.00180	0.00225	0.00225
50 Trin196735	0.00226	0.00180	0.00180	0.00180	0.00180	0.00180	0.00226	0.00226
51 pall207682	0.00271	0.00317	0.00317	0.00317	0.00317	0.00317	0.00271	0.00271
52 pallWR8626	0.00591	0.00637	0.00637	0.00637	0.00637	0.00637	0.00591	0.00591
53 wagn010616	0.07859	0.07917	0.07917	0.07917	0.07917	0.07917	0.07859	0.07859
54 wagn012969	0.07983	0.08041	0.08041	0.08041	0.08041	0.08041	0.07983	0.07983
55 wagn177351	0.07974	0.08032	0.08032	0.08032	0.08032	0.08032	0.07974	0.07974
56 podicUC243	0.13826	0.13899	0.13899	0.13899	0.13899	0.13899	0.13826	0.13826
57 podicUC244	0.13761	0.13834	0.13834	0.13834	0.13834	0.13834	0.13761	0.13761
58 podicUC278	0.14460	0.14535	0.14535	0.14535	0.14535	0.14535	0.14460	0.14460
59 podic53124	0.14119	0.14194	0.14194	0.14194	0.14194	0.14194	0.14119	0.14119
60 chaq186524	0.17386	0.17424	0.17424	0.17424	0.17424	0.17424	0.17386	0.17386
61 knuds13244	0.22738	0.22830	0.22830	0.22830	0.22830	0.22830	0.22738	0.22738

	17	18	19	20	21	22	23	24
17 Gren197007	-							
18 Gren197008	0.00000	-						
19 Gren197017	0.00045	0.00045	-					
20 Gren197044	0.00000	0.00000	0.00045	-				
21 StVn056421	0.00045	0.00045	0.00000	0.00045	-			
22 StVn056490	0.00045	0.00045	0.00000	0.00045	0.00000	-		
23 StVn056561	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000	-	
24 StVn056562	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000	0.00000	-
25 StVn056612	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000	0.00000	0.00000
26 StVn196895	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000	0.00000	0.00000
27 StVn196896	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045	0.00045	0.00045
28 StVn196897	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045	0.00045	0.00045
29 StVn196898	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000	0.00000	0.00000
30 StVn196899	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045	0.00045	0.00045
31 StVn196900	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045	0.00045	0.00045
32 Tobo186597	0.00271	0.00271	0.00225	0.00271	0.00225	0.00225	0.00225	0.00225
33 Trin196726	0.00135	0.00135	0.00180	0.00135	0.00180	0.00180	0.00180	0.00180
34 Trin196727	0.00135	0.00135	0.00180	0.00135	0.00180	0.00180	0.00180	0.00180
35 Trin196728	0.00180	0.00180	0.00225	0.00180	0.00225	0.00225	0.00225	0.00225
36 Trin196729	0.00180	0.00180	0.00135	0.00180	0.00135	0.00135	0.00135	0.00135
37 Trin196730	0.00135	0.00135	0.00180	0.00135	0.00180	0.00180	0.00180	0.00180
38 Trin196731	0.00135	0.00135	0.00180	0.00135	0.00180	0.00180	0.00180	0.00180
39 Trin196732	0.00135	0.00135	0.00180	0.00135	0.00180	0.00180	0.00180	0.00180
40 Trin196733	0.00135	0.00135	0.00180	0.00135	0.00180	0.00180	0.00180	0.00180
41 Trin196734	0.00135	0.00135	0.00180	0.00135	0.00180	0.00180	0.00180	0.00180
42 Trin196888	0.00135	0.00135	0.00180	0.00135	0.00180	0.00180	0.00180	0.00180
43 StVn056613	0.00045	0.00045	0.00000	0.00045	0.00000	0.00000	0.00000	0.00000
44 StVn196894	0.00000	0.00000	0.00045	0.00000	0.00045	0.00045	0.00045	0.00045
45 Trin175424	0.00271	0.00271	0.00225	0.00271	0.00225	0.00225	0.00225	0.00225
46 Tobo186596	0.00271	0.00271	0.00225	0.00271	0.00225	0.00225	0.00225	0.00225
47 Trin196886	0.00316	0.00316	0.00271	0.00316	0.00271	0.00271	0.00271	0.00271
48 Trin175410	0.00135	0.00135	0.00180	0.00135	0.00180	0.00180	0.00180	0.00180
49 Trin175620	0.00180	0.00180	0.00225	0.00180	0.00225	0.00225	0.00225	0.00225
50 Trin196735	0.00180	0.00180	0.00226	0.00180	0.00226	0.00226	0.00226	0.00226
51 pall207682	0.00317	0.00317	0.00271	0.00317	0.00271	0.00271	0.00271	0.00271
52 pallWR8626	0.00637	0.00637	0.00591	0.00637	0.00591	0.00591	0.00591	0.00591
53 wagn010616	0.07917	0.07917	0.07859	0.07917	0.07859	0.07859	0.07859	0.07859
54 wagn012969	0.08041	0.08041	0.07983	0.08041	0.07983	0.07983	0.07983	0.07983
55 wagn177351	0.08032	0.08032	0.07974	0.08032	0.07974	0.07974	0.07974	0.07974
56 podicUC243	0.13899	0.13899	0.13826	0.13899	0.13826	0.13826	0.13826	0.13826
57 podicUC244	0.13834	0.13834	0.13761	0.13834	0.13761	0.13761	0.13761	0.13761
58 podicUC278	0.14535	0.14535	0.14460	0.14535	0.14460	0.14460	0.14460	0.14460
59 podic53124	0.14194	0.14194	0.14119	0.14194	0.14119	0.14119	0.14119	0.14119
60 chaq186524	0.17424	0.17424	0.17386	0.17424	0.17386	0.17386	0.17386	0.17386
61 knuds13244	0.22830	0.22830	0.22738	0.22830	0.22738	0.22738	0.22738	0.22738

	25	26	27	28	29	30	31	32
25 StVn056612	-							
26 StVn196895	0.00000	-						
27 StVn196896	0.00045	0.00045	-					
28 StVn196897	0.00045	0.00045	0.00000	-				
29 StVn196898	0.00000	0.00000	0.00045	0.00045	-			
30 StVn196899	0.00045	0.00045	0.00000	0.00000	0.00045	-		
31 StVn196900	0.00045	0.00045	0.00000	0.00000	0.00045	0.00000	-	
32 Tobo186597	0.00225	0.00225	0.00271	0.00271	0.00225	0.00271	0.00271	-
33 Trin196726	0.00180	0.00180	0.00135	0.00135	0.00180	0.00135	0.00135	0.00135
34 Trin196727	0.00180	0.00180	0.00135	0.00135	0.00180	0.00135	0.00135	0.00135
35 Trin196728	0.00225	0.00225	0.00180	0.00180	0.00225	0.00180	0.00180	0.00180
36 Trin196729	0.00135	0.00135	0.00180	0.00180	0.00135	0.00180	0.00180	0.00090
37 Trin196730	0.00180	0.00180	0.00135	0.00135	0.00180	0.00135	0.00135	0.00135
38 Trin196731	0.00180	0.00180	0.00135	0.00135	0.00180	0.00135	0.00135	0.00135
39 Trin196732	0.00180	0.00180	0.00135	0.00135	0.00180	0.00135	0.00135	0.00135
40 Trin196733	0.00180	0.00180	0.00135	0.00135	0.00180	0.00135	0.00135	0.00135
41 Trin196734	0.00180	0.00180	0.00135	0.00135	0.00180	0.00135	0.00135	0.00135
42 Trin196888	0.00180	0.00180	0.00135	0.00135	0.00180	0.00135	0.00135	0.00135
43 StVn056613	0.00000	0.00000	0.00045	0.00045	0.00000	0.00045	0.00045	0.00225
44 StVn196894	0.00045	0.00045	0.00000	0.00000	0.00045	0.00000	0.00000	0.00271
45 Trin175424	0.00225	0.00225	0.00271	0.00271	0.00225	0.00271	0.00271	0.00180
46 Tobo186596	0.00225	0.00225	0.00271	0.00271	0.00225	0.00271	0.00271	0.00000
47 Trin196886	0.00271	0.00271	0.00316	0.00316	0.00271	0.00316	0.00316	0.00226
48 Trin175410	0.00180	0.00180	0.00135	0.00135	0.00180	0.00135	0.00135	0.00135
49 Trin175620	0.00225	0.00225	0.00180	0.00180	0.00225	0.00180	0.00180	0.00180
50 Trin196735	0.00226	0.00226	0.00180	0.00180	0.00226	0.00180	0.00180	0.00180
51 pall207682	0.00271	0.00271	0.00317	0.00317	0.00271	0.00317	0.00317	0.00225
52 pallWR8626	0.00591	0.00591	0.00637	0.00637	0.00591	0.00637	0.00637	0.00546
53 wagn010616	0.07859	0.07859	0.07917	0.07917	0.07859	0.07917	0.07917	0.07788
54 wagn012969	0.07983	0.07983	0.08041	0.08041	0.07983	0.08041	0.08041	0.07912
55 wagn177351	0.07974	0.07974	0.08032	0.08032	0.07974	0.08032	0.08032	0.07903
56 podicUC243	0.13826	0.13826	0.13899	0.13899	0.13826	0.13899	0.13899	0.13694
57 podicUC244	0.13761	0.13761	0.13834	0.13834	0.13761	0.13834	0.13834	0.13630
58 podicUC278	0.14460	0.14460	0.14535	0.14535	0.14460	0.14535	0.14535	0.14327
59 podic53124	0.14119	0.14119	0.14194	0.14194	0.14119	0.14194	0.14194	0.13986
60 chaq186524	0.17386	0.17386	0.17424	0.17424	0.17386	0.17424	0.17424	0.17211
61 knuds13244	0.22738	0.22738	0.22830	0.22830	0.22738	0.22830	0.22830	0.22517

	33	34	35	36	37	38	39	40
33 Trin196726	-							
34 Trin196727	0.00000	-						
35 Trin196728	0.00045	0.00045	-					
36 Trin196729	0.00045	0.00045	0.00090	-				
37 Trin196730	0.00000	0.00000	0.00045	0.00045	-			
38 Trin196731	0.00000	0.00000	0.00045	0.00045	0.00000	-		
39 Trin196732	0.00000	0.00000	0.00045	0.00045	0.00000	0.00000	-	
40 Trin196733	0.00000	0.00000	0.00045	0.00045	0.00000	0.00000	0.00000	-
41 Trin196734	0.00000	0.00000	0.00045	0.00045	0.00000	0.00000	0.00000	0.00000
42 Trin196888	0.00000	0.00000	0.00045	0.00045	0.00000	0.00000	0.00000	0.00000
43 StVn056613	0.00180	0.00180	0.00225	0.00135	0.00180	0.00180	0.00180	0.00180
44 StVn196894	0.00135	0.00135	0.00180	0.00180	0.00135	0.00135	0.00135	0.00135
45 Trin175424	0.00135	0.00135	0.00180	0.00090	0.00135	0.00135	0.00135	0.00135
46 Tobol186596	0.00135	0.00135	0.00180	0.00090	0.00135	0.00135	0.00135	0.00135
47 Trin196886	0.00180	0.00180	0.00226	0.00135	0.00180	0.00180	0.00180	0.00180
48 Trin175410	0.00000	0.00000	0.00045	0.00045	0.00000	0.00000	0.00000	0.00000
49 Trin175620	0.00045	0.00045	0.00090	0.00090	0.00045	0.00045	0.00045	0.00045
50 Trin196735	0.00045	0.00045	0.00090	0.00090	0.00045	0.00045	0.00045	0.00045
51 pall207682	0.00180	0.00180	0.00225	0.00135	0.00180	0.00180	0.00180	0.00180
52 pallWR8626	0.00500	0.00500	0.00546	0.00454	0.00500	0.00500	0.00500	0.00500
53 wagn010616	0.07714	0.07714	0.07790	0.07656	0.07714	0.07714	0.07714	0.07714
54 wagn012969	0.07836	0.07836	0.07914	0.07778	0.07836	0.07836	0.07836	0.07836
55 wagn177351	0.07828	0.07828	0.07905	0.07770	0.07828	0.07828	0.07828	0.07828
56 podicUC243	0.13735	0.13735	0.13730	0.13662	0.13735	0.13735	0.13735	0.13735
57 podicUC244	0.13670	0.13670	0.13666	0.13597	0.13670	0.13670	0.13670	0.13670
58 podicUC278	0.14366	0.14366	0.14363	0.14292	0.14366	0.14366	0.14366	0.14366
59 podic53124	0.14026	0.14026	0.14019	0.13951	0.14026	0.14026	0.14026	0.14026
60 chaq186524	0.17434	0.17434	0.17551	0.17396	0.17434	0.17434	0.17434	0.17434
61 knuds13244	0.22565	0.22565	0.22440	0.22474	0.22565	0.22565	0.22565	0.22565

	41	42	43	44	45	46	47	48
41 Trin196734	-							
42 Trin196888	0.00000	-						
43 StVn056613	0.00180	0.00180	-					
44 StVn196894	0.00135	0.00135	0.00045	-				
45 Trin175424	0.00135	0.00135	0.00225	0.00271	-			
46 Tobol186596	0.00135	0.00135	0.00225	0.00271	0.00180	-		
47 Trin196886	0.00180	0.00180	0.00271	0.00316	0.00226	0.00226	-	
48 Trin175410	0.00000	0.00000	0.00180	0.00135	0.00135	0.00135	0.00181	-
49 Trin175620	0.00045	0.00045	0.00225	0.00180	0.00180	0.00180	0.00226	0.00045
50 Trin196735	0.00045	0.00045	0.00226	0.00180	0.00180	0.00180	0.00226	0.00045
51 pall207682	0.00180	0.00180	0.00271	0.00317	0.00225	0.00225	0.00271	0.00180
52 pallWR8626	0.00500	0.00500	0.00591	0.00637	0.00546	0.00546	0.00595	0.00500
53 wagn010616	0.07714	0.07714	0.07859	0.07917	0.07788	0.07788	0.07732	0.07715
54 wagn012969	0.07836	0.07836	0.07983	0.08041	0.07911	0.07912	0.07856	0.07838
55 wagn177351	0.07828	0.07828	0.07974	0.08032	0.07903	0.07903	0.07847	0.07830
56 podicUC243	0.13735	0.13735	0.13826	0.13899	0.13630	0.13694	0.13564	0.13675
57 podicUC244	0.13670	0.13670	0.13761	0.13834	0.13566	0.13630	0.13500	0.13611
58 podicUC278	0.14366	0.14366	0.14460	0.14535	0.14259	0.14327	0.14192	0.14305
59 podic53124	0.14026	0.14026	0.14119	0.14194	0.13918	0.13986	0.13852	0.13966
60 chaq186524	0.17434	0.17434	0.17386	0.17424	0.17351	0.17211	0.17389	0.17441
61 knuds13244	0.22565	0.22565	0.22738	0.22830	0.22683	0.22517	0.22475	0.22576



	49	50	51	52	53	54	55	56
49 Trin175620	-							
50 Trin196735	0.00090	-						
51 pall207682	0.00225	0.00226	-					
52 pallWR8626	0.00547	0.00546	0.00409	-				
53 wagn010616	0.07639	0.07773	0.07658	0.07412	-			
54 wagn012969	0.07916	0.07896	0.07780	0.07686	0.01651	-		
55 wagn177351	0.07754	0.07888	0.07773	0.07525	0.00226	0.01600	-	
56 podicUC243	0.13579	0.13811	0.13585	0.13517	0.12881	0.12854	0.13078	-
57 podicUC244	0.13514	0.13747	0.13521	0.13453	0.12816	0.12791	0.13013	0.00045
58 podicUC278	0.14206	0.14339	0.14213	0.14143	0.13226	0.13405	0.13431	0.00781
59 podic53124	0.13868	0.13997	0.13873	0.13803	0.13389	0.13507	0.13594	0.01491
60 chaq186524	0.17326	0.17353	0.17501	0.16933	0.16905	0.17899	0.17048	0.15571
61 knuds13244	0.22452	0.22661	0.22664	0.22271	0.23176	0.23572	0.23535	0.21656

	57	58	59	60	61
57 podicUC244	-				
58 podicUC278	0.00734	-			
59 podic53124	0.01444	0.01636	-		
60 chaq186524	0.15503	0.15858	0.15744	-	
61 knuds13244	0.21573	0.22178	0.21699	0.17204	-

Table XII. Absolute base differences among samples for the combined dataset.

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
1 Gren006881	-														
2 Gren006882	0	-													
3 Gren006883	0	0	-												
4 Gren006939	0	0	0	-											
5 Gren196977	1	1	1	1	-										
6 Gren196978	0	0	0	0	1	-									
7 Gren196979	1	1	1	1	0	1	-								
8 Gren196980	1	1	1	1	0	1	0	-							
9 Gren196999	0	0	0	0	1	0	1	1	-						
10 Gren197000	1	1	1	1	0	1	0	0	1	-					
11 Gren197001	1	1	1	1	0	1	0	0	1	0	-				
12 Gren197002	1	1	1	1	0	1	0	0	1	0	0	-			
13 Gren197003	1	1	1	1	0	1	0	0	1	0	0	0	-		
14 Gren197004	1	1	1	1	0	1	0	0	1	0	0	0	0	-	
15 Gren197005	0	0	0	0	1	0	1	1	0	1	1	1	1	1	-
16 Gren197006	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
17 Gren197007	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
18 Gren197008	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
19 Gren197017	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
20 Gren197044	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
21 StVn056421	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
22 StVn056490	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
23 StVn056561	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
24 StVn056562	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
25 StVn056612	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
26 StVn196895	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
27 StVn196896	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
28 StVn196897	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
29 StVn196898	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
30 StVn196899	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
31 StVn196900	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
32 Tobol86597	5	5	5	5	6	5	6	6	5	6	6	6	6	6	5
33 Trin196726	4	4	4	4	3	4	3	3	4	3	3	3	3	3	4
34 Trin196727	4	4	4	4	3	4	3	3	4	3	3	3	3	3	4
35 Trin196728	5	5	5	5	4	5	4	4	5	4	4	4	4	4	5
36 Trin196729	3	3	3	3	4	3	4	4	3	4	4	4	4	4	3
37 Trin196730	4	4	4	4	3	4	3	3	4	3	3	3	3	3	4
38 Trin196731	4	4	4	4	3	4	3	3	4	3	3	3	3	3	4
39 Trin196732	4	4	4	4	3	4	3	3	4	3	3	3	3	3	4
40 Trin196733	4	4	4	4	3	4	3	3	4	3	3	3	3	3	4
41 Trin196734	4	4	4	4	3	4	3	3	4	3	3	3	3	3	4
42 Trin196888	4	4	4	4	3	4	3	3	4	3	3	3	3	3	4
43 StVn056613	0	0	0	0	1	0	1	1	0	1	1	1	1	1	0
44 StVn196894	1	1	1	1	0	1	0	0	1	0	0	0	0	0	1
45 Trin175424	5	5	5	5	6	5	6	6	5	6	6	6	6	6	5
46 Tobol86596	5	5	5	5	6	5	6	6	5	6	6	6	6	6	5
47 Trin196886	6	6	6	6	7	6	7	7	6	7	7	7	7	7	6
48 Trin175410	4	4	4	4	3	4	3	3	4	3	3	3	3	3	4
49 Trin175620	5	5	5	5	4	5	4	4	5	4	4	4	4	4	5
50 Trin196735	5	5	5	5	4	5	4	4	5	4	4	4	4	4	5
51 pall207682	6	6	6	6	7	6	7	7	6	7	7	7	7	7	6
52 pallWR8626	13	13	13	13	14	13	14	14	13	14	14	14	14	14	13
53 wagn010616	142	142	142	142	143	142	143	143	142	143	143	143	143	143	142
54 wagn012969	143	143	143	143	144	143	144	144	143	144	144	144	144	144	143
55 wagn177351	144	144	144	144	145	144	145	145	144	145	145	145	145	145	144
56 podicUC243	228	228	228	228	229	228	229	229	228	229	229	229	229	229	228
57 podicUC244	227	227	227	227	228	227	228	228	227	228	228	228	228	228	227
58 podicUC278	235	235	235	235	236	235	236	236	235	236	236	236	236	236	235
59 podic53124	231	231	231	231	232	231	232	232	231	232	232	232	232	232	231
60 chaq186524	269	269	269	269	269	269	269	269	269	269	269	269	269	269	269
61 knuds13244	335	335	335	335	336	335	336	336	335	336	336	336	336	336	335

	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30
16 Gren197006	-														
17 Gren197007	1	-													
18 Gren197008	1	0	-												
19 Gren197017	0	1	1	-											
20 Gren197044	1	0	0	1	-										
21 StVn056421	0	1	1	0	1	-									
22 StVn056490	0	1	1	0	1	0	-								
23 StVn056561	0	1	1	0	1	0	0	-							
24 StVn056562	0	1	1	0	1	0	0	0	-						
25 StVn056612	0	1	1	0	1	0	0	0	0	-					
26 StVn196895	0	1	1	0	1	0	0	0	0	0	-				
27 StVn196896	1	0	0	1	0	1	1	1	1	1	1	-			
28 StVn196897	1	0	0	1	0	1	1	1	1	1	1	0	-		
29 StVn196898	0	1	1	0	1	0	0	0	0	0	0	1	1	-	
30 StVn196899	1	0	0	1	0	1	1	1	1	1	1	1	0	1	-
31 StVn196900	1	0	0	1	0	1	1	1	1	1	1	1	0	0	1
32 Tobol86597	5	6	6	5	6	5	5	5	5	5	5	6	6	5	6
33 Trin196726	4	3	3	4	3	4	4	4	4	4	4	4	3	3	4
34 Trin196727	4	3	3	4	3	4	4	4	4	4	4	4	3	3	4
35 Trin196728	5	4	4	5	4	5	5	5	5	5	5	5	4	4	5
36 Trin196729	3	4	4	3	4	3	3	3	3	3	3	4	4	3	4
37 Trin196730	4	3	3	4	3	4	4	4	4	4	4	4	3	3	4
38 Trin196731	4	3	3	4	3	4	4	4	4	4	4	4	3	3	4
39 Trin196732	4	3	3	4	3	4	4	4	4	4	4	4	3	3	4
40 Trin196733	4	3	3	4	3	4	4	4	4	4	4	4	3	3	4
41 Trin196734	4	3	3	4	3	4	4	4	4	4	4	4	3	3	4
42 Trin196888	4	3	3	4	3	4	4	4	4	4	4	4	3	3	4
43 StVn056613	0	1	1	0	1	0	0	0	0	0	0	1	1	0	1
44 StVn196894	1	0	0	1	0	1	1	1	1	1	1	1	0	0	1
45 Trin175424	5	6	6	5	6	5	5	5	5	5	5	6	6	5	6
46 Tobol86596	5	6	6	5	6	5	5	5	5	5	5	6	6	5	6
47 Trin196886	6	7	7	6	7	6	6	6	6	6	6	7	7	6	7
48 Trin175410	4	3	3	4	3	4	4	4	4	4	4	4	3	3	4
49 Trin175620	5	4	4	5	4	5	5	5	5	5	5	5	4	4	5
50 Trin196735	5	4	4	5	4	5	5	5	5	5	5	5	4	4	5
51 pall1207682	6	7	7	6	7	6	6	6	6	6	6	7	7	6	7
52 pallWR8626	13	14	14	13	14	13	13	13	13	13	13	14	14	13	14
53 wagn010616	142	143	143	142	143	142	142	142	142	142	142	143	143	142	143
54 wagn012969	143	144	144	143	144	143	143	143	143	143	143	144	144	143	144
55 wagn177351	144	145	145	144	145	144	144	144	144	144	144	145	145	144	145
56 podicUC243	228	229	229	228	229	228	228	228	228	228	228	229	229	228	229
57 podicUC244	227	228	228	227	228	227	227	227	227	227	227	228	228	227	228
58 podicUC278	235	236	236	235	236	235	235	235	235	235	235	236	236	235	236
59 podic53124	231	232	232	231	232	231	231	231	231	231	231	232	232	231	232
60 chaq186524	269	269	269	269	269	269	269	269	269	269	269	269	269	269	269
61 knuds13244	335	336	336	335	336	335	335	335	335	335	335	336	336	335	336

	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45
31 StVn196900	-														
32 Tobol86597	6	-													
33 Trin196726	3	3	-												
34 Trin196727	3	3	0	-											
35 Trin196728	4	4	1	1	-										
36 Trin196729	4	2	1	1	2	-									
37 Trin196730	3	3	0	0	1	1	-								
38 Trin196731	3	3	0	0	1	1	0	-							
39 Trin196732	3	3	0	0	1	1	0	0	-						
40 Trin196733	3	3	0	0	1	1	0	0	0	-					
41 Trin196734	3	3	0	0	1	1	0	0	0	0	-				
42 Trin196888	3	3	0	0	1	1	0	0	0	0	0	-			
43 StVn056613	1	5	4	4	5	3	4	4	4	4	4	4	-		
44 StVn196894	0	6	3	3	4	4	3	3	3	3	3	3	1	-	
45 Trin175424	6	4	3	3	4	2	3	3	3	3	3	3	5	6	-
46 Tobol86596	6	0	3	3	4	2	3	3	3	3	3	3	5	6	4
47 Trin196886	7	5	4	4	5	3	4	4	4	4	4	4	6	7	5
48 Trin175410	3	3	0	0	1	1	0	0	0	0	0	0	4	3	3
49 Trin175620	4	4	1	1	2	2	1	1	1	1	1	1	5	4	4
50 Trin196735	4	4	1	1	2	2	1	1	1	1	1	1	5	4	4
51 pall207682	7	5	4	4	5	3	4	4	4	4	4	4	6	7	5
52 pallWR8626	14	12	11	11	12	10	11	11	11	11	11	11	13	14	12
53 wagn010616	143	141	140	140	141	139	140	140	140	140	140	140	142	143	141
54 wagn012969	144	142	141	141	142	140	141	141	141	141	141	141	143	144	142
55 wagn177351	145	143	142	142	143	141	142	142	142	142	142	142	144	145	143
56 podicUC243	229	226	227	227	227	226	227	227	227	227	227	227	228	229	226
57 podicUC244	228	225	226	226	226	225	226	226	226	226	226	226	227	228	225
58 podicUC278	236	233	234	234	234	233	234	234	234	234	234	234	235	236	233
59 podic53124	232	229	230	230	230	229	230	230	230	230	230	230	231	232	229
60 chaq186524	269	267	269	269	270	269	269	269	269	269	269	269	269	269	269
61 knuds13244	336	332	333	333	332	332	333	333	333	333	333	333	335	336	334

	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60
46 Tobol86596	-														
47 Trin196886	5	-													
48 Trin175410	3	4	-												
49 Trin175620	4	5	1	-											
50 Trin196735	4	5	1	2	-										
51 pall207682	5	6	4	5	5	-									
52 pallWR8626	12	13	11	12	12	9	-								
53 wagn010616	141	140	140	139	141	139	135	-							
54 wagn012969	142	141	141	142	142	140	138	35	-						
55 wagn177351	143	142	142	141	143	141	137	5	34	-					
56 podicUC243	226	225	226	225	228	225	224	215	213	217	-				
57 podicUC244	225	224	225	224	227	224	223	214	212	216	1	-			
58 podicUC278	233	232	233	232	234	232	231	218	219	220	17	16	-		
59 podic53124	229	228	229	228	230	228	227	220	220	222	32	31	35	-	
60 chaq186524	267	269	269	268	268	270	264	263	273	264	249	248	253	252	-
61 knuds13244	332	332	333	332	334	334	330	336	340	339	325	324	330	326	274

61 knuds13244 -

## Appendix I

Museum catalogue numbers, field numbers, and locality data for the samples utilized in this study. Museum abbreviations: BWMC = Bobby Witcher Memorial Collection, Avila University, LSUZ = Louisiana State University, Museum of Natural Science, QCAZ = Quito Catholic Zoology Museum, USNM = United States National Museum (USA).

Species	Museum Number	Field Number	Locality
<i>L. chaquensis</i>	USNM 319708	USFS 186524	Tucumán; ca. 40 km SE San Miguel de Tucumán, Argentina.
<i>L. knudseni</i>	QCAZ 13244		Provincia de Orellana; Parque Nacional Yasuní, Ecuador.
<i>L. pallidirostris</i>	USNM 535774	USFS 207682	Northwest District; Baramita, Guyana.
<i>L. pallidirostris</i>	USNM 302408	WRH 8626	Roraima; Igarapé Cocal, Brazil.
<i>L. podicipinus</i>		UC 243	Mato Grosso do Sol; Estância Mimosa; Mun de Bonito, Brazil.
<i>L. podicipinus</i>		UC 244	Mato Grosso do Sol; Estância Mimosa; Mun de Bonito, Brazil.
<i>L. podicipinus</i>		UC 278	São Paulo; Chácara Renascer; Bauru, Brazil.
<i>L. podicipinus</i>	USNM 053124	USFS 303207	São Paulo; Fazenda Jatai, Brazil.
<i>L. validus</i>	BWMC 06881		St. Andrew; Spring Gardens Estate, Grenada.
<i>L. validus</i>	BWMC 06882		St. Andrew; Birch Grove, Grenada.
<i>L. validus</i>	BWMC 06883		St. Andrew; Birch Grove, Grenada.
<i>L. validus</i>	BWMC 06939		St. George; Beausejour, Grenada.
<i>L. validus</i>	USNM 314793	USFS 196977	St. George; Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314794	USFS 196978	St. George; Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314795	USFS 196979	St. George; Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314796	USFS 196980	St. George; Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314798	USFS 197044	St. George; Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314813	USFS 196999	St. George; Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314814	USFS 197000	St. George; inland from Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314815	USFS 197001	St. George; inland from Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314816	USFS 197002	St. George; inland from Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314817	USFS 197003	St. George; inland from Grand Anse Bay, Grenada.

Species	Museum Number	Field Number	Locality
<i>L. validus</i>	USNM 314818	USFS 197004	St. George; inland from Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314819	USFS 197005	St. George; inland from Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314820	USFS 197006	St. George; inland from Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314821	USFS 197007	St. George; inland from Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314822	USFS 197008	St. George; inland from Grand Anse Bay, Grenada.
<i>L. validus</i>	USNM 314831	USFS 197017	St. George; inland from Grand Anse Beach, Grenada.
<i>L. validus</i>	USNM 314512	USFS 56421	St. Andrew; near Vermont, St. Vincent.
<i>L. validus</i>	USNM 314513	USFS 56490	St. George; Arnos Vale, St. Vincent.
<i>L. validus</i>	USNM 314514	USFS 56612	St. George; Arnos Vale, St. Vincent.
<i>L. validus</i>	USNM 314515	USFS 56613	St. George; Arnos Vale, St. Vincent.
<i>L. validus</i>	USNM 314718	USFS 196894	St. George; Arnos Vale, St. Vincent.
<i>L. validus</i>	USNM 314719	USFS 196895	St. George; Arnos Vale, St. Vincent.
<i>L. validus</i>	USNM 314720	USFS 196896	St. George; Arnos Vale, St. Vincent.
<i>L. validus</i>	USNM 314721	USFS 196897	St. George; Arnos Vale, St. Vincent.
<i>L. validus</i>	USNM 314722	USFS 196898	St. George; Arnos Vale, St. Vincent.
<i>L. validus</i>	USNM 314723	USFS 196899	St. George; Arnos Vale, St. Vincent.
<i>L. validus</i>	USNM 314714	USFS 196900	St. George; Arnos Vale, St. Vincent.
<i>L. validus</i>	USNM 314516	USFS 56561	St. George; Rose Cottage, St. Vincent.
<i>L. validus</i>	USNM 314517	USFS 56562	St. George; Rose Cottage, St. Vincent.
<i>L. validus</i>	USNM 523940	USFS 186596	St. Paul; Delaford, Louis d'Or River, Tobago.
<i>L. validus</i>	USNM 523941	USFS 186597	St. Paul; Dealford, Louis d'Or River, Tobago.
<i>L. validus</i>	USNM 286948	USFS 175410	St. George; Simla Research Station, Trinidad.
<i>L. validus</i>	USNM 286959	USFS 175424	St. George; north of Simla Research Station, Trinidad.
<i>L. validus</i>	USNM 306105	USFS 175620	St. George; near Brasso Seco Village, Trinidad.
<i>L. validus</i>	USNM 314671	USFS 196886	St. George; west of Carapo, Trinidad.
<i>L. validus</i>	USNM 314672	USFS 196888	St. George; west of Carapo, Trinidad.
<i>L. validus</i>	USNM 314627	USFS 196726	St. Patrick; near Chatham Beach, Trinidad.
<i>L. validus</i>	USNM 314628	USFS 196727	St. Patrick; near Chatham Beach, Trinidad.
<i>L. validus</i>	USNM 314629	USFS 196728	St. Patrick; near Chatham Beach, Trinidad.

Species	Museum Number	Field Number	Locality
<i>L. validus</i>	USNM 314630	USFS 196729	St. Patrick; near Chatham Beach, Trinidad.
<i>L. validus</i>	USNM 314631	USFS 196730	St. Patrick; near Chatham Beach, Trinidad.
<i>L. validus</i>	USNM 314632	USFS 196731	St. Patrick; near Chatham Beach, Trinidad.
<i>L. validus</i>	USNM 314633	USFS 196732	St. Patrick; near Chatham Beach, Trinidad.
<i>L. validus</i>	USNM 314634	USFS 196733	St. Patrick; near Chatham Beach, Trinidad.
<i>L. validus</i>	USNM 314635	USFS 196734	St. Patrick; near Chatham Beach, Trinidad.
<i>L. validus</i>	USNM 314636	USFS 196735	St. Patrick; near Chatham Beach, Trinidad.
<i>L. wagneri</i>	LSUMZ H-13653	JPC 12969	Acre; ca. 5km N Porto Walter, inland from the Rio Jurua, Brazil.
<i>L. wagneri</i>	LSUMZ H-12885	JPC 10616	Sucumbios; Estación, Científica near Cuyabeno, Ecuador.
<i>L. wagneri</i>	USNM 320988	USFS 177351	Pastaza; Coca, 1km ENE of Tiguino, Ecuador.

## Appendix II

Alignment of 12S sequences. Shaded regions denoting stem positions in the secondary structure model are numbered arbitrarily. Numbering below the alignment begins at the first 12S base position of the *Xenopus laevis* mitochondrial genome. Insertions in the *X. laevis* sequence are represented by diagonal shading, and were excluded from analyses.

	1			2			1'			3			3a			4																																			
Gren006881	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G				
Gren006882	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren006883	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren006939	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren196977	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren196978	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren196979	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren196980	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren196999	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren197000	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren197001	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren197002	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren197003	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren197004	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren197005	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren197006	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren197007	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren197008	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren197017	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Gren197044	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
StVn056421	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
StVn056490	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
StVn056561	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
StVn056562	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
StVn056612	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
StVn196895	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
StVn196896	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
StVn196897	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
StVn196898	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
StVn196899	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
StVn196900	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Tobo186597	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Trin196726	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Trin196727	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Trin196728	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Trin196729	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Trin196730	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Trin196731	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Trin196732	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Trin196733	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Trin196734	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Trin196888	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
StVn056613	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
StVn196894	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A	C	C	T	T	A	A	A	A	T	C	A	A	C	T	T	T	A	C	T	T	A	C	T	T	A	C	T	T	A	C	A	C	A	T	G
Trin175424	C	A	A	A	G	G	T	T	G	G	T	C	C	T	G	A																																			







































	33	33'	22a'	22'	34
Gren006881	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren006882	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren006883	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren006939	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren196977	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren196978	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren196979	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren196980	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren196999	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren197000	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren197001	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren197002	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren197003	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren197004	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren197005	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren197006	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren197007	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren197008	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren197017	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Gren197044	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
StVn056421	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
StVn056490	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
StVn056561	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
StVn056562	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
StVn056612	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
StVn196895	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
StVn196896	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
StVn196897	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
StVn196898	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
StVn196899	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
StVn196900	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Tobol186597	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin196726	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin196727	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin196728	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin196729	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin196730	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin196731	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin196732	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin196733	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin196734	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin196888	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
StVn056613	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
StVn196894	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin175424	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Tobol186596	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin196886	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin175410	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin175620	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
Trin196735	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
pall1207682	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
pallWR8626	AAACAGAGT	GTCCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
wagn010616	AAACAGAGT	GTTCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
wagn012969	AAACAGAGT	GTTCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
wagn177351	AAACAGAGT	GTTCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
podicUC243	AAACAGAGT	GTTCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
podicUC244	AAACAGAGT	GTTCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
podicUC278	AAACAGAGT	GTTCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
podic53124	AAACAGAGT	GTTCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
chaq186524	AAACAGAGA	GTTCTTTT	TAAACCC	GGCAC	TGGGATGTGTACACACCG
knuds13244	GACATAGT	GCTCTTTT	TAAATCC	GGCAC	TGGGATGTGTACACACCG
Xenopuslav	AAACAGAGA	GTTCTCTT	TAAAAAC	GGCCC	TGGGATGTGTACACACCG

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	17'	18	18'
Gren006881	CCAG	CCTAAAAT	TACTCCAATGCA
Gren006882	CCAG	CCTAAAAT	TACTCCAATGCA
Gren006883	CCAG	CCTAAAAT	TACTCCAATGCA
Gren006939	CCAG	CCTAAAAT	TACTCCAATGCA
Gren196977	CCAG	CCTAAAAT	TACTCCAATGCA
Gren196978	CCAG	CCTAAAAT	TACTCCAATGCA
Gren196979	CCAG	CCTAAAAT	TACTCCAATGCA
Gren196980	CCAG	CCTAAAAT	TACTCCAATGCA
Gren196999	CCAG	CCTAAAAT	TACTCCAATGCA
Gren197000	CCAG	CCTAAAAT	TACTCCAATGCA
Gren197001	CCAG	CCTAAAAT	TACTCCAATGCA
Gren197002	CCAG	CCTAAAAT	TACTCCAATGCA
Gren197003	CCAG	CCTAAAAT	TACTCCAATGCA
Gren197004	CCAG	CCTAAAAT	TACTCCAATGCA
Gren197005	CCAG	CCTAAAAT	TACTCCAATGCA
Gren197006	CCAG	CCTAAAAT	TACTCCAATGCA
Gren197007	CCAG	CCTAAAAT	TACTCCAATGCA
Gren197008	CCAG	CCTAAAAT	TACTCCAATGCA
Gren197017	CCAG	CCTAAAAT	TACTCCAATGCA
Gren197044	CCAG	CCTAAAAT	TACTCCAATGCA
StVn056421	CCAG	CCTAAAAT	TACTCCAATGCA
StVn056490	CCAG	CCTAAAAT	TACTCCAATGCA
StVn056561	CCAG	CCTAAAAT	TACTCCAATGCA
StVn056562	CCAG	CCTAAAAT	TACTCCAATGCA
StVn056612	CCAG	CCTAAAAT	TACTCCAATGCA
StVn196895	CCAG	CCTAAAAT	TACTCCAATGCA
StVn196896	CCAG	CCTAAAAT	TACTCCAATGCA
StVn196897	CCAG	CCTAAAAT	TACTCCAATGCA
StVn196898	CCAG	CCTAAAAT	TACTCCAATGCA
StVn196899	CCAG	CCTAAAAT	TACTCCAATGCA
StVn196900	CCAG	CCTAAAAT	TACTCCAATGCA
Tobo186597	CCAG	CCTAAAAT	TACTCCAATGCA
Trin196726	CCAG	CCTAAAAT	TACTCCAATGCA
Trin196727	CCAG	CCTAAAAT	TACTCCAATGCA
Trin196728	CCAG	CCTAAAAT	TACTCCAATGCA
Trin196729	CCAG	CCTAAAAT	TACTCCAATGCA
Trin196730	CCAG	CCTAAAAT	TACTCCAATGCA
Trin196731	CCAG	CCTAAAAT	TACTCCAATGCA
Trin196732	CCAG	CCTAAAAT	TACTCCAATGCA
Trin196733	CCAG	CCTAAAAT	TACTCCAATGCA
Trin196734	CCAG	CCTAAAAT	TACTCCAATGCA
Trin196888	CCAG	CCTAAAAT	TACTCCAATGCA
StVn056613	CCAG	CCTAAAAT	TACTCCAATGCA
StVn196894	CCAG	CCTAAAAT	TACTCCAATGCA
Trin175424	CCAG	CCTAAAAT	TACTCCAATGCA
Tobo186596	CCAG	CCTAAAAT	TACTCCAATGCA
Trin196886	CCAG	CCTAAAAT	TACTCCAATGCA
Trin175410	CCAG	CCTAAAAT	TACTCCAATGCA
Trin175620	CCAG	CCTAAAAT	TACTCCAATGCA
Trin196735	CCAG	CCTAAAAT	TACTCCAATGCA
pall1207682	CCAG	CCTAAAAT	TACTCCAATGCA
pallWR8626	CCAG	CCTAAAAT	TACTCCAATGCA
wagn010616	CCAA	CCTAAAAT	TACTTCTATATA
wagn012969	CCAA	CCTAAAAT	TACTTCTATATA
wagn177351	CCAA	CCTAAAAT	TACTTCTATATA
podicUC243	CCAG	CCTAAAAT	TATTTCTATATT
podicUC244	CCAG	CCTAAAAT	TATTTCTATATT
podicUC278	CCAG	CCTAAAAT	TATTTCTATATT
podic53124	CCAG	CCTAAAAT	TATTTCTATATT
chaq186524	CCAA	CCTAAAAC	TACTTAAATTA
knuds13244	CCAA	CCTAAAAC	TACTTATATCC
Xenopuslav	TCTA	CCCTAAATA	TAGATTTTAAAG









	26	26'	9'
Gren006881	T T C T A T	A A C C C C T	A T A G A A A C T
Gren006882	T T C T A T	A A C C C C T	A T A G A A A C T
Gren006883	T T C T A T	A A C C C C T	A T A G A A A C T
Gren006939	T T C T A T	A A C C C C T	A T A G A A A C T
Gren196977	T T C T A T	A A C C C C T	A T A G A A A C T
Gren196978	T T C T A T	A A C C C C T	A T A G A A A C T
Gren196979	T T C T A T	A A C C C C T	A T A G A A A C T
Gren196980	T T C T A T	A A C C C C T	A T A G A A A C T
Gren196999	T T C T A T	A A C C C C T	A T A G A A A C T
Gren197000	T T C T A T	A A C C C C T	A T A G A A A C T
Gren197001	T T C T A T	A A C C C C T	A T A G A A A C T
Gren197002	T T C T A T	A A C C C C T	A T A G A A A C T
Gren197003	T T C T A T	A A C C C C T	A T A G A A A C T
Gren197004	T T C T A T	A A C C C C T	A T A G A A A C T
Gren197005	T T C T A T	A A C C C C T	A T A G A A A C T
Gren197006	T T C T A T	A A C C C C T	A T A G A A A C T
Gren197007	T T C T A T	A A C C C C T	A T A G A A A C T
Gren197008	T T C T A T	A A C C C C T	A T A G A A A C T
Gren197017	T T C T A T	A A C C C C T	A T A G A A A C T
Gren197044	T T C T A T	A A C C C C T	A T A G A A A C T
StVn056421	T T C T A T	A A C C C C T	A T A G A A A C T
StVn056490	T T C T A T	A A C C C C T	A T A G A A A C T
StVn056561	T T C T A T	A A C C C C T	A T A G A A A C T
StVn056562	T T C T A T	A A C C C C T	A T A G A A A C T
StVn056612	T T C T A T	A A C C C C T	A T A G A A A C T
StVn196895	T T C T A T	A A C C C C T	A T A G A A A C T
StVn196896	T T C T A T	A A C C C C T	A T A G A A A C T
StVn196897	T T C T A T	A A C C C C T	A T A G A A A C T
StVn196898	T T C T A T	A A C C C C T	A T A G A A A C T
StVn196899	T T C T A T	A A C C C C T	A T A G A A A C T
StVn196900	T T C T A T	A A C C C C T	A T A G A A A C T
Tobol186597	T T C T A T	A A C C C - T	A T A G A A A C T
Trin196726	T T C T A T	A A C C C - T	A T A G A A A C T
Trin196727	T T C T A T	A A C C C - T	A T A G A A A C T
Trin196728	T T C T A T	A A C C C - T	A T A G A A A C T
Trin196729	T T C T A T	A A C C C - T	A T A G A A A C T
Trin196730	T T C T A T	A A C C C - T	A T A G A A A C T
Trin196731	T T C T A T	A A C C C - T	A T A G A A A C T
Trin196732	T T C T A T	A A C C C - T	A T A G A A A C T
Trin196733	T T C T A T	A A C C C - T	A T A G A A A C T
Trin196734	T T C T A T	A A C C C - T	A T A G A A A C T
Trin196888	T T C T A T	A A C C C - T	A T A G A A A C T
StVn056613	T T C T A T	A A C C C C T	A T A G A A A C T
StVn196894	T T C T A T	A A C C C C T	A T A G A A A C T
Trin175424	T T C T A T	A A C C C - T	A T A G A A A C T
Tobol186596	T T C T A T	A A C C C - T	A T A G A A A C T
Trin196886	T T C T A T	A A C C C - T	A T A G A A A C T
Trin175410	T T C T A T	A A C C C - T	A T A G A A A C T
Trin175620	T T C T A T	A A C C C - T	A T A G A A A C T
Trin196735	T T C T A T	A A C C C - T	A T A G A A A C T
pall1207682	T T C T A T	A A C C C - T	A T A G A A A C T
pallWR8626	T T C T A T	A A C C C - T	A T A G A A A C T
wagn010616	T T C T A T	A A A A T T T	A T A G A A A C T
wagn012969	T T C T A T	A A A A T T T	A T A G A A A C T
wagn177351	T T C T A T	A A A A T T T	A T A G A A A C T
podicUC243	T T C T A T	A T A A C - T	A T A G A A A C T
podicUC244	T T C T A T	A T A A C - T	A T A G A A A C T
podicUC278	T T C T A T	A T A A C - T	A T A G A A A C T
podic53124	T T C T A T	A T A A C - T	A T A G A A A C T
chaq186524	T T C T A T	A A T A C - T	A T A G A A A C T
knuds13244	T T C T A T	A T A T C - T	A T A G A A A C T
Xenopuslav	T T C T A T	A A A C - - T	A T A G A A A C T































	54	54'	55	56	56'	57
Gren006881	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren006882	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren006883	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren006939	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren196977	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren196978	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren196979	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren196980	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren196999	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren197000	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren197001	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren197002	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren197003	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren197004	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren197005	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren197006	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren197007	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren197008	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren197017	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Gren197044	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
StVn056421	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
StVn056490	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
StVn056561	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
StVn056562	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
StVn056612	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
StVn196895	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
StVn196896	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
StVn196897	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
StVn196898	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
StVn196899	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
StVn196900	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Tobol86597	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin196726	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin196727	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin196728	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin196729	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin196730	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin196731	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin196732	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin196733	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin196734	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin196888	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
StVn056613	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
StVn196894	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin175424	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Tobol86596	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin196886	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin175410	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin175620	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
Trin196735	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
pall207682	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
pallWR8626	AGCAAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	CTAAATTC
wagn010616	AGTAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	-TAAATTC
wagn012969	AGTAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	-TAAATTC
wagn177351	AGTAAAAAATAAAC	CTCCACAGTGA	-ATGGGAC	CTCT	TCCC	-TAAATTC
podicUC243	AGCAAAAAAATAAAC	CTCCACAGTGA	AAAAAGGAC	CTT	TCCC	CTAAATTC
podicUC244	AGCAAAAAAATAAAC	CTCCACAGTGA	AAAAAGGAC	CTT	TCCC	CTAAATTC
podicUC278	AGCAAAAAAATAAAC	CTCCACAGTGA	AAAAAGGAC	CTT	TCCC	CTAAATTC
podic53124	AGCAAAAAAATAAAC	CTCCACAGTGA	AAAAAGGAC	CTT	TCCC	CTAAATTC
chaq186524	AGCAAAAAAATAAAC	CTCCGCAGTGA	-ATGGGAC	CTTAT	TCCC	-TAAATTC
knuds13244	AGTAAAAAATAAAC	CTCCGCAGTGA	-ACAGGGAC	CTCT	TCCC	-TAAATTC
Xenopuslav	AGATAATAATAATC	CTCCTTGA	AGA	-ATAGGGC	CTATCA	CCCT-TTCA

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	65'
Gren006881	A C C
Gren006882	A C C
Gren006883	A C C
Gren006939	A C C
Gren196977	A C C
Gren196978	A C C
Gren196979	A C C
Gren196980	A C C
Gren196999	A C C
Gren197000	A C C
Gren197001	A C C
Gren197002	A C C
Gren197003	A C C
Gren197004	A C C
Gren197005	A C C
Gren197006	A C C
Gren197007	A C C
Gren197008	A C C
Gren197017	A C C
Gren197044	A C C
StVn056421	A C C
StVn056490	A C C
StVn056561	A C C
StVn056562	A C C
StVn056612	A C C
StVn196895	A C C
StVn196896	A C C
StVn196897	A C C
StVn196898	A C C
StVn196899	A C C
StVn196900	A C C
Tobol86597	A C C
Trin196726	A C C
Trin196727	A C C
Trin196728	A C C
Trin196729	A C C
Trin196730	A C C
Trin196731	A C C
Trin196732	A C C
Trin196733	A C C
Trin196734	A C C
Trin196888	A C C
StVn056613	A C C
StVn196894	A C C
Trin175424	A C C
Tobol86596	A C C
Trin196886	A C C
Trin175410	A C C
Trin175620	A C C
Trin196735	A C C
pall207682	A C C
pallWR8626	A C C
wagn010616	A C C
wagn012969	A C C
wagn177351	A C C
podicUC243	A C C
podicUC244	A C C
podicUC278	A C C
podic53124	A C C
chaq186524	A C C
knuds13244	A C C
Xenopuslav	G C C

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## Vita

Kenneth Christopher Yanek was born on 4 June 1977 in Rockville Centre, New York, USA. He graduated from Adelphi University, Garden City, New York, where he completed the requirements for the degree of Bachelor of Science in Biology in May, 2001. He completed the requirements for the degree of Master of Science in Biology in July 2004.