



Identification of *Leishmania* spp. and a report of novel species in Thailand

*Saovanee Leelayoova, Suradej Siripattanapipong,
Theerayudh Sukmee, Tawee Naaglor,
Jeerapun Worapong and Mathirut Mungthin*



Diagnosis of Leishmaniasis

- Traditional diagnostic methods
- Serological diagnosis
- Molecular diagnosis
- Species identification

Traditional diagnostic methods

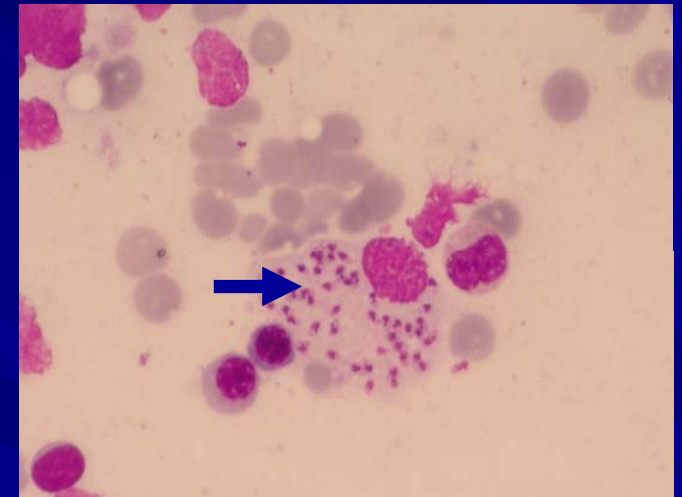


Microscopic examination

- Examination of intracellular amastigotes from Giemsa-stained lesion biopsy smears (CL), or lymph node, bone marrow aspiration (VL)
- remains the gold standard in leishmania diagnosis
- High specificity

Disadvantages

- Operator dependent
- low sensitivity
- procedures for demonstration of the parasite are invasive



Intracellular amastigotes



Traditional diagnostic methods

■ Culture

- Growth of promastigotes in days to weeks

■ Sensitivity of 50-70%*



* Weigle KA et al, Am J Trop Med Hyg 1987; 36: 489-496

Cultivation of Promastigotes (1)



■ Undefined medium

- NNN medium with Locke's solution overlays
- NNN medium with medium-199 as a liquid phase

■ Partially or Semi-defined liquid media

- Grace's medium
- Schneider's Drosophila medium,
- M199

All require a supplement of 10-20% heat inactivated bovine serum or calf serum

Cultivation of Promastigotes (2)



■ Define liquid media

(serum-free media)

- Enriched Synthetic Medium (ESM)
- Medium 199+
- RPMI+

Culture conditions : 26° C or room temperature, adequate aeration



Promastigote, flagellate form



Serological Diagnosis

- Indirect fluorescence antibody (IFA)
- ELISA

Not suitable for field conditions

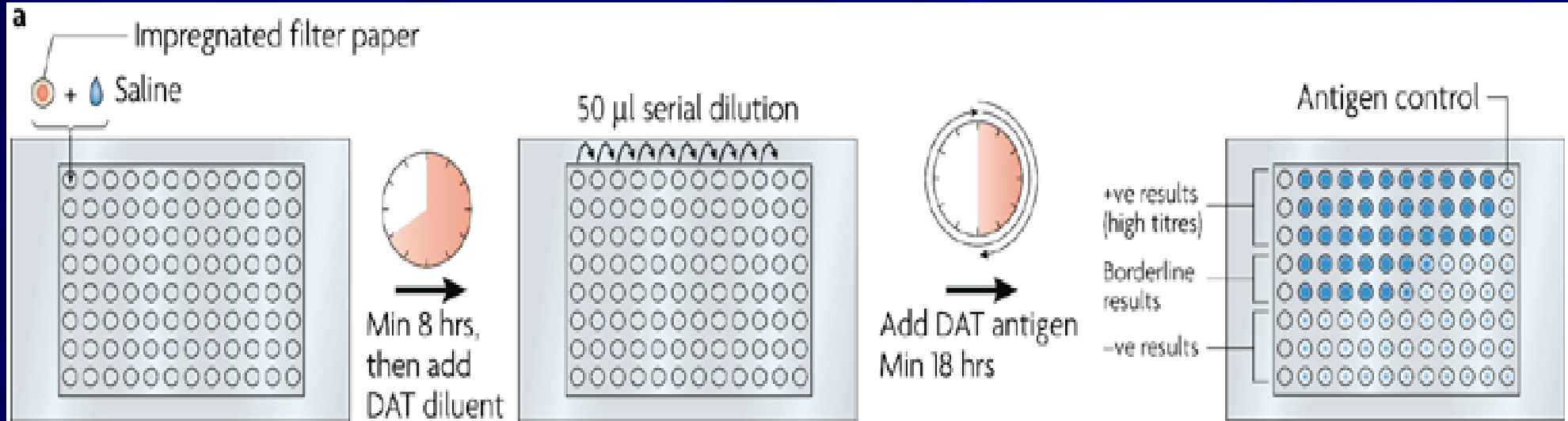
Field use

- Direct Agglutination Test (DAT)
- rK39
- *Fast Agglutination Screening Test (FAST)



DAT and rK39 dipstick

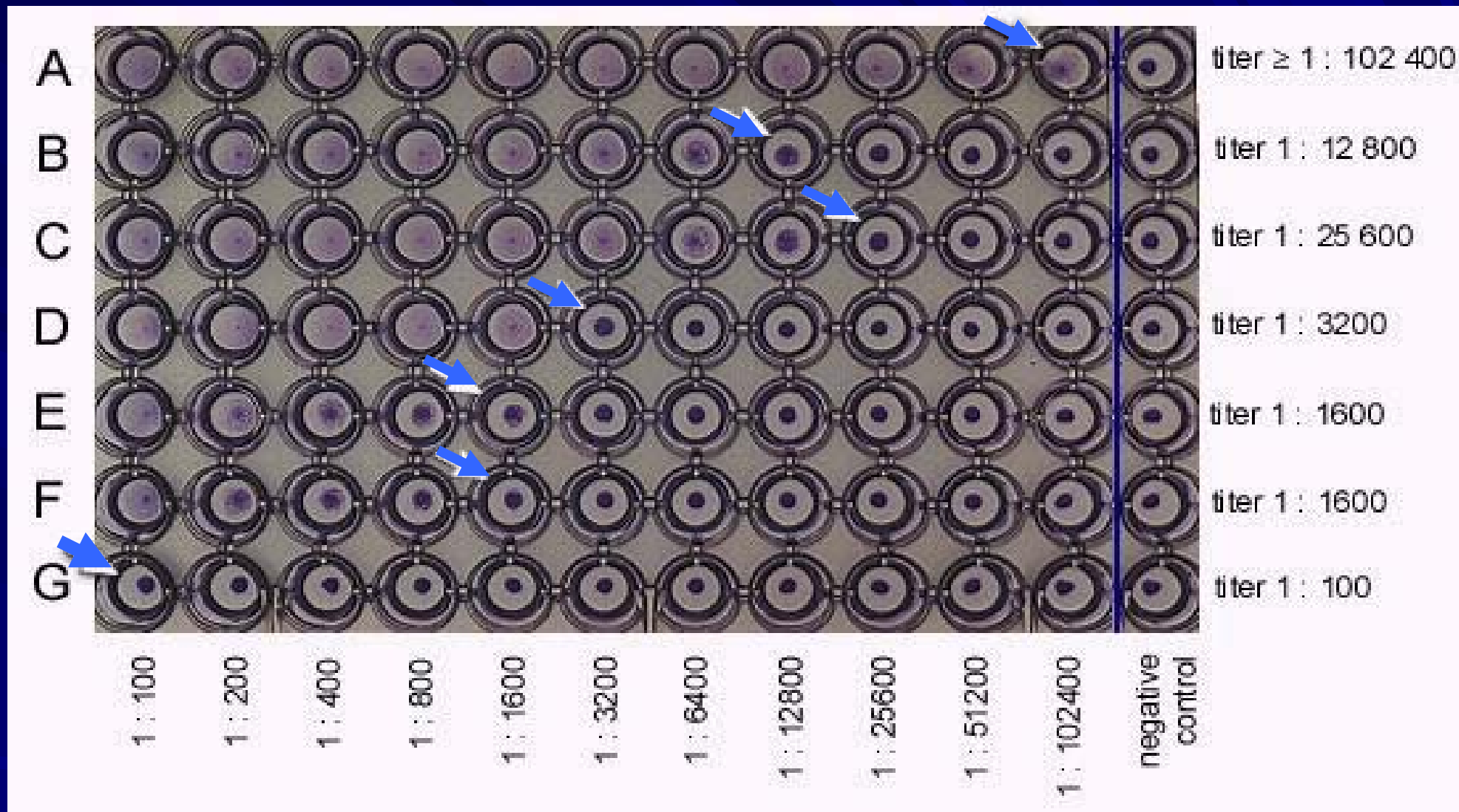
- Easy to use
- Require minimal technological expertise or laboratory set up
- Ideal for both field and laboratory use



Direct Agglutination Test (DAT)

- A semi-quantitative test
- using dilutions of patient's serum and stained killed *L. donovani* promastigotes
- Agglutination is visible after 18 hours with the naked eyes
- Develop for field use

DAT



Limitation of use: Agglutinating antibodies may persist for a long time after treatment. The DAT does not discriminate between current and past infection.

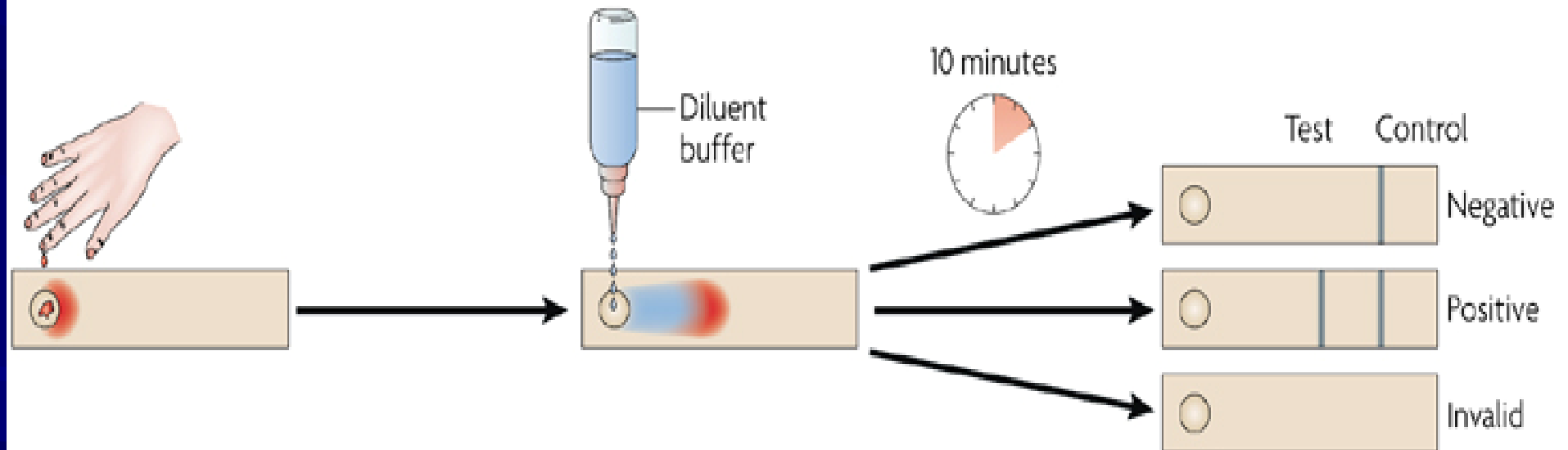


rK39

- Recombinant kinesin-related protein of 39 kDa called rK39
- a part of a kinesin-related protein in *L. chagasi* which is conserved within *L. donovani* complex
- antibody in cutaneous and mucocutaneous leishmaniasis can not be detected

rK39 : immunochromatographic test strip

- developed to dipstick





Evaluation of DAT and rK39 dipstick for visceral leishmaniasis

- The Meta analysis of 30 studies (1986-2004)*
- Similar diagnostic performance
good → excellent
- Seem comparable
- **Sensitivities**
 - DAT 94.8% (95%CI, 92.7-96.4%)
 - rK39 dipstick 93.9% (95%CI, 87.7-97.1%)
- **Specificities**
 - DAT 85.9% (95%CI, 93.9-98.7%)
 - rK39 dipstick 90.6% (95%CI, 66.8-97.9%)

*Chappuis F et.al., *BMJ* 2006;333: 723



rKRP42

- a promising new antigen
- part of an *L. donovani* kinesin-related protein
- rKRP42 having 39 more amino acids than rK39.

Test for the sensitivity and specificity

sensitivity	<u>94.6%</u> (70 /74 VL samples)
specificity	99.3% (148/149 negative samples)

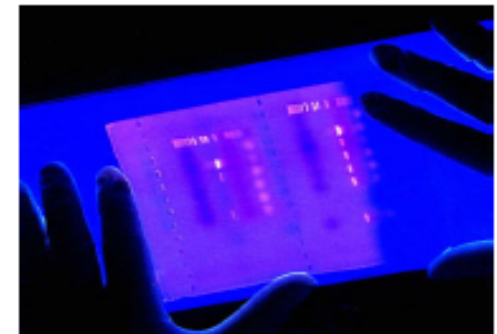
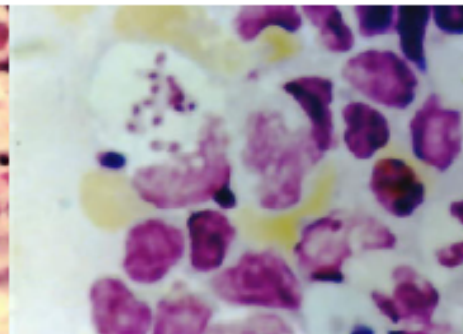
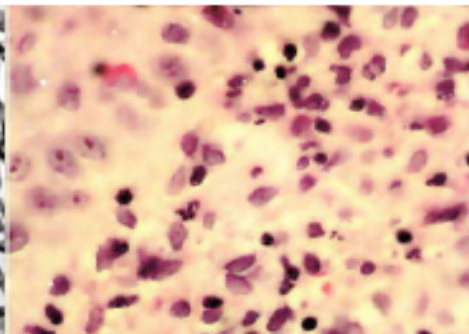
- sensitivity of rK39 dipstick test was 93.2% (69 /74 VL patients).

Leishmaniasis Diagnosis: Comparison of different conventional methods and PCR/Hybridization

n=156



n=23



Culture

HE

TP

PCR/Hyb

LCL

44%

50%

35%

94%

ML

10%

26%

0%

70%

*Dept. of Tropical Medicine, Instituto Oswaldo Cruz, Fundação Oswaldo Cruz
Ministry of Health, Rio de Janeiro, Brasil*

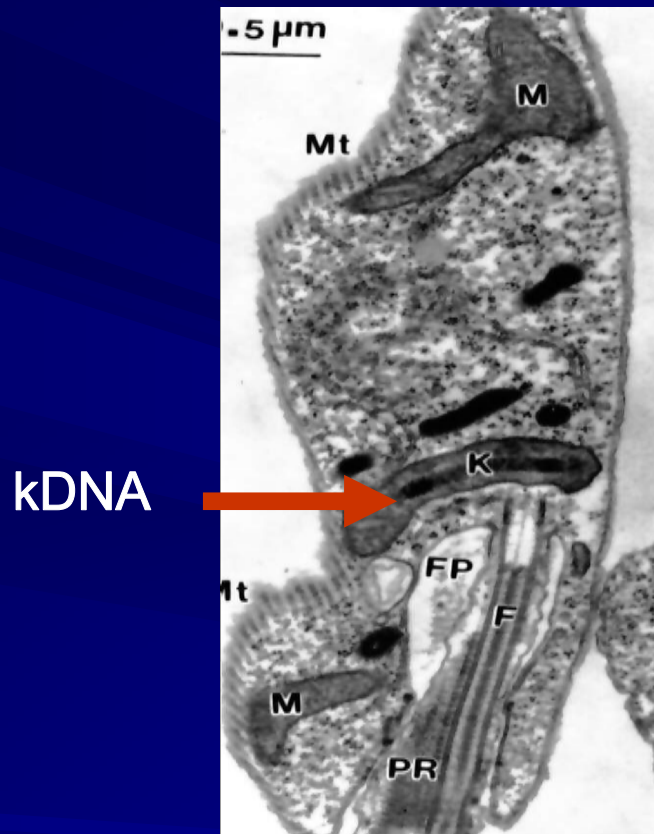


PCR and Leishmaniasis

- Over 400 articles published on PCR and leishmaniasis since 1989
- **High-copy-number target genes are chosen**
 - rRNA genes
 - ITS1 region of rRNA genes
 - Kinetoplast DNA minicircles
 - Mini-exon genes
- **Other genes**
 - Gp63 genes
 - hsp70 genes
 - Cysteine proteinase genes



kDNA minicircle gene



- An excellent target for a sensitive and rapid detection method
- present at thousands of copies per cell
- show the highest sensitivity by PCR



PCR (kDNA gene) & Hybridization

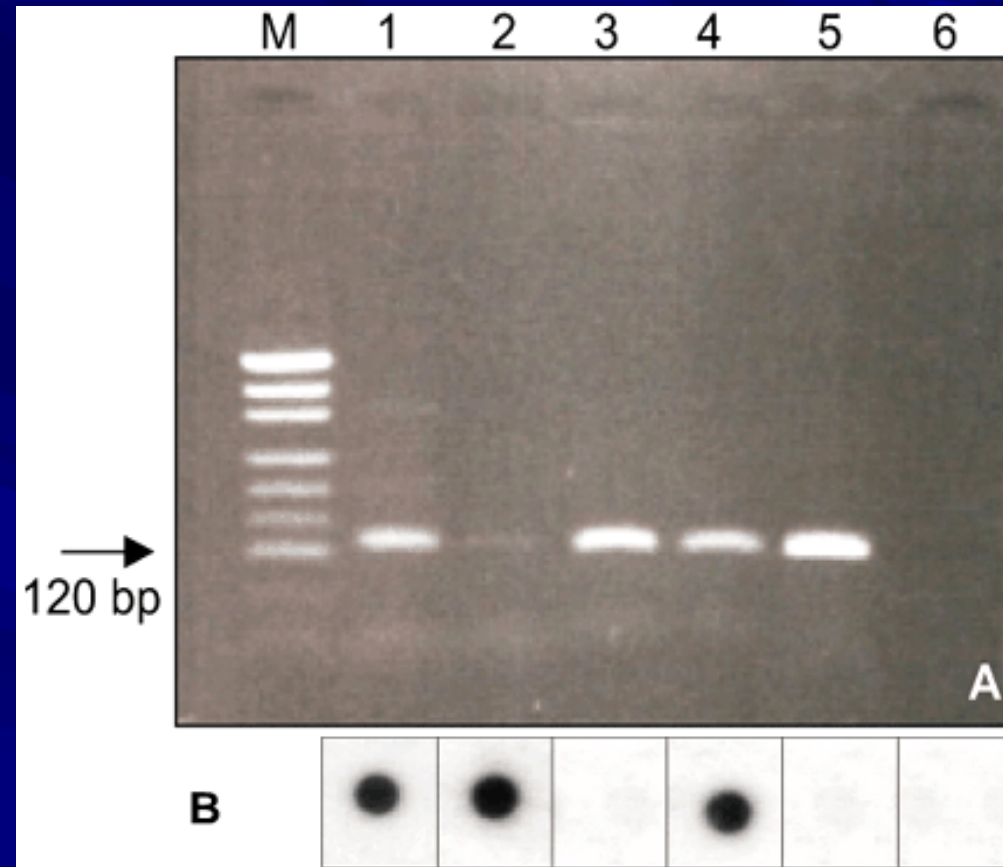
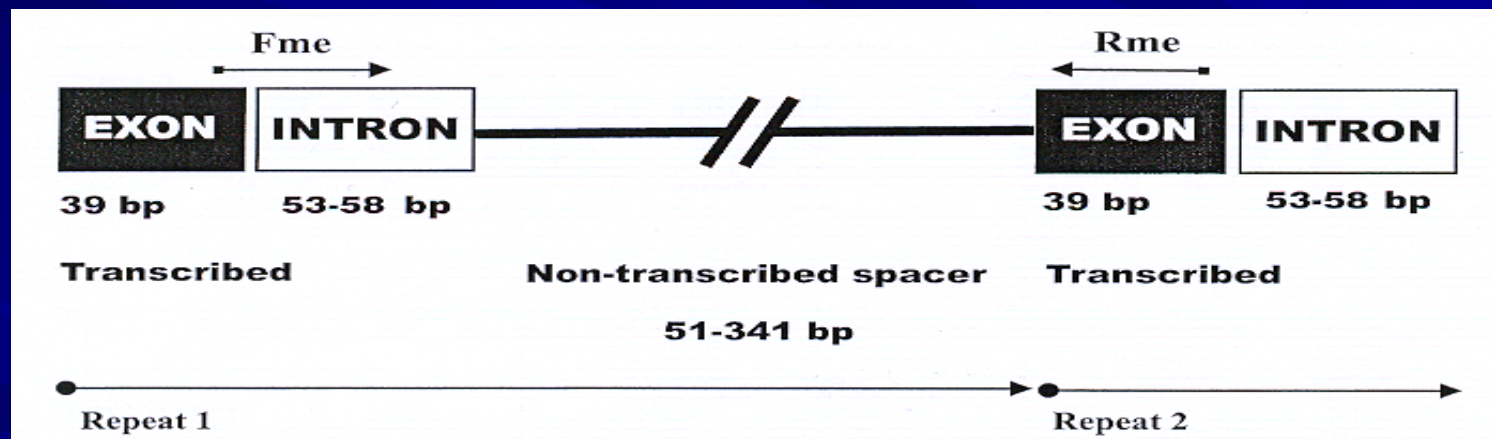


Fig. 2 - A. PCR products obtained with primers of the genus *Leishmania* visualized after 2% agarose gel electrophoresis stained with ethidium bromide. MW 50base-pair DNA ladder; lanes 1 and 2: bone marrow and blood from patient, lane 03: DNA of *L. chagasi*, lane 4: DNA of *L. braziliensis*, lane 5: DNA of *L. amazonensis* and lane 6: negative control. **B.** Hybridization with a cloned minicircle from *L. braziliensis*.



Mini-exon (spliced leader) gene repeat

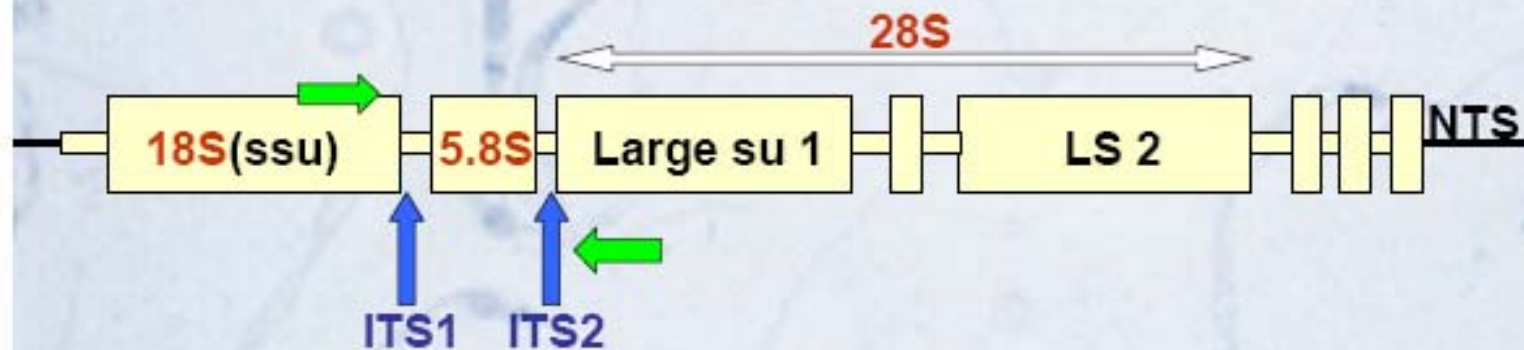
- Involve in the transsplicing process of nuclear mRNA
- Present 100-200 tandemly repeated copies per nuclear genome
- Sequence variation of non-transcribed spacer was observed between individual species



Marfurt J et. al., J Clin microbiol 2003; 41: 3147-3153.

rRNA Gene : 16S-23S / Internal Transcribed Spacers

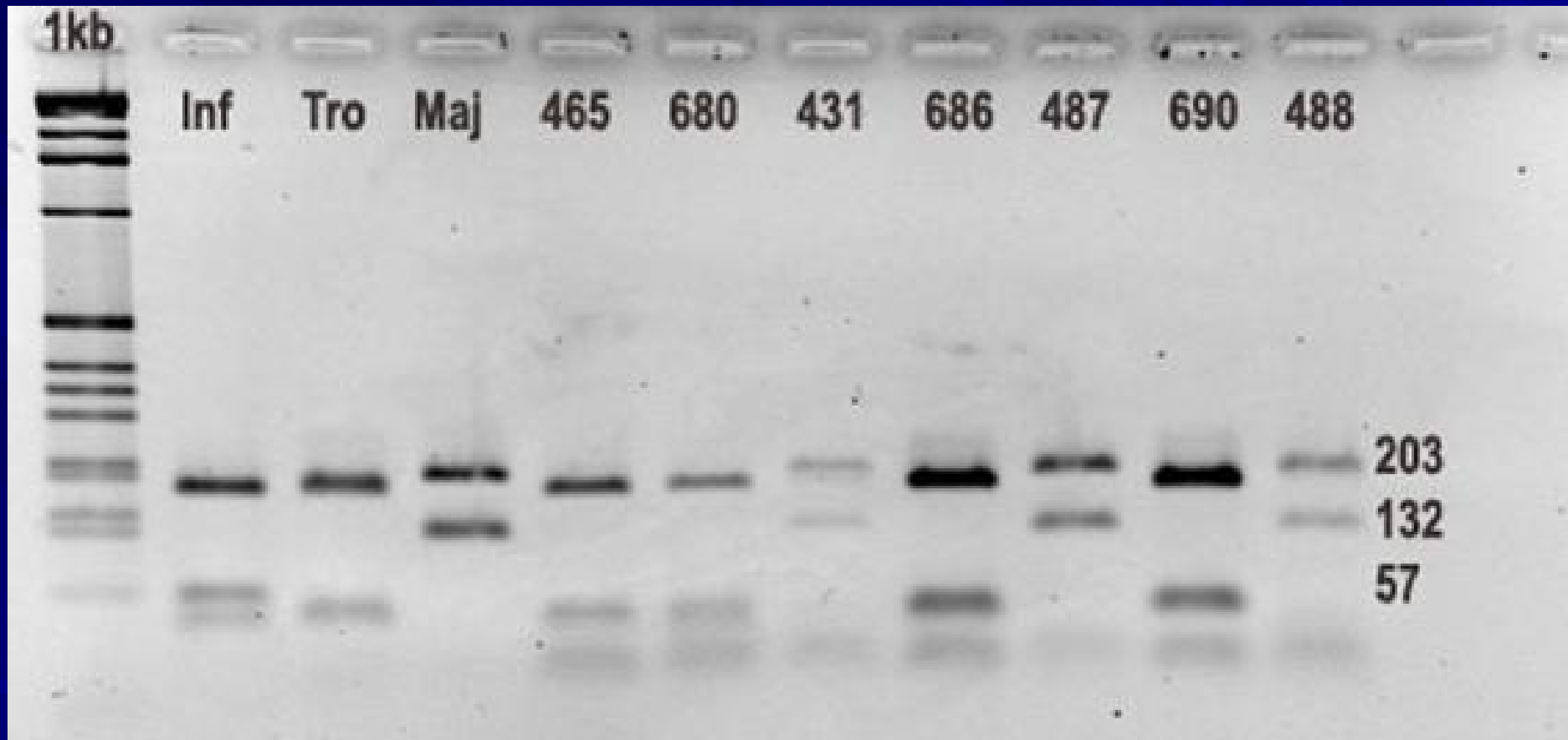
r RNA locus contains between the small sub-unit 18S and the large subunit, 2 variable sequences ITS 1 and ITS 2



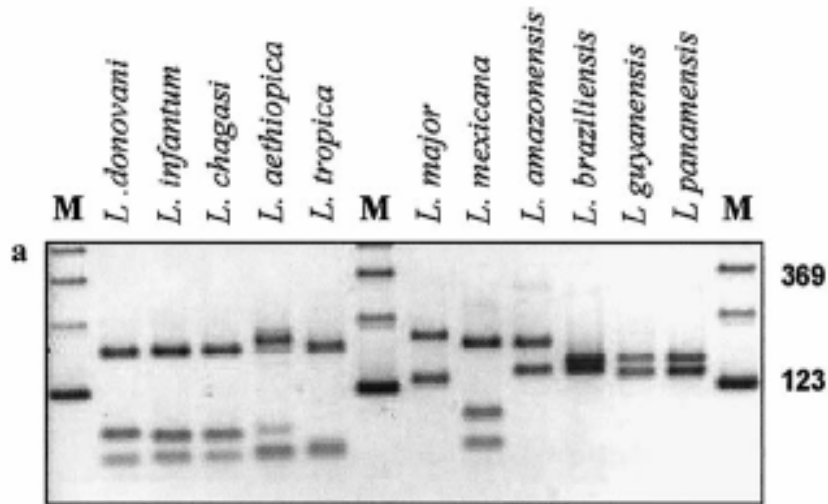
It was possible to amplify this region with 2 primers located between the Ssu and the Lsu



PCR-RFLP (ITS1)

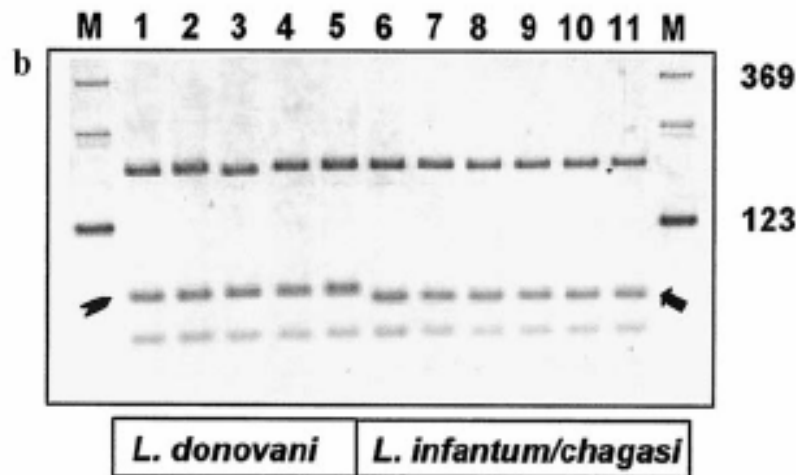


Using 1 restriction enzyme, *HaeIII*



ITS1 region of SSU rRNA genes

- RFLP using enzyme *HaeIII*



* Schonian G, Nasereddin A, Dinse N, Schweynoch C, Schallig HDFH, Presber W, et al. *Diagn Microbiol Infect Dis.* 2003;47:349-58.

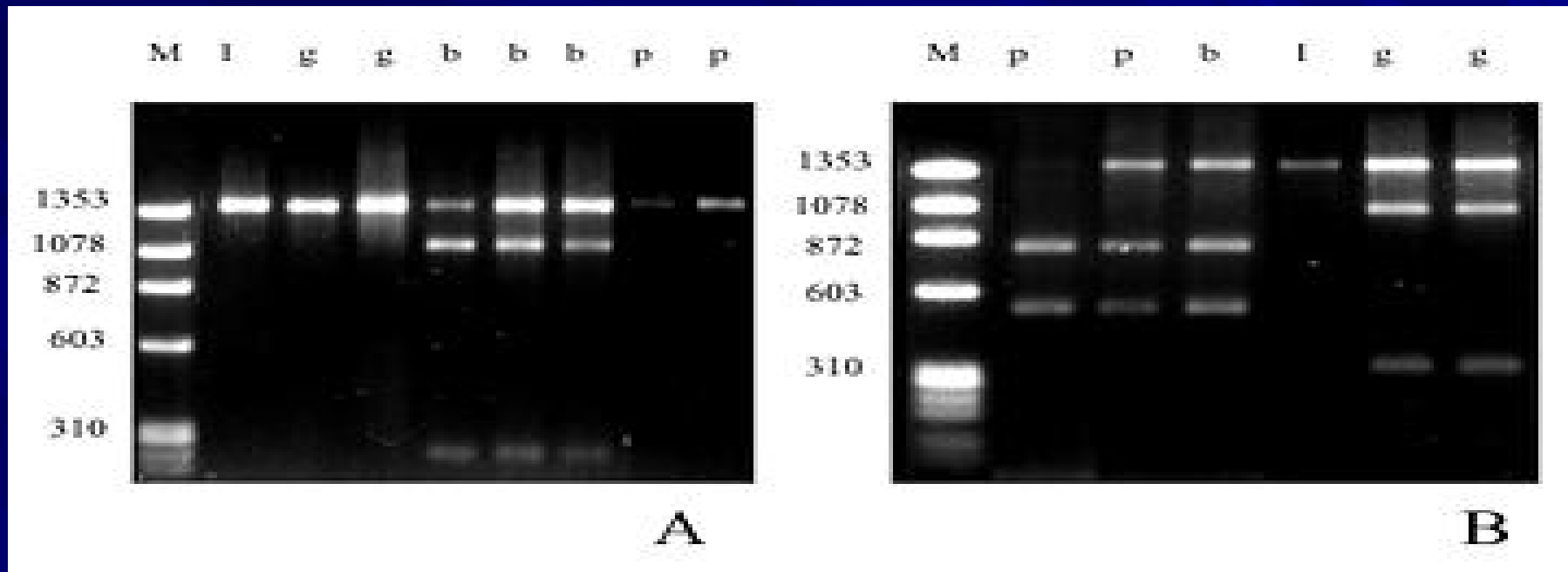


Gp63

- Promastigote surface protease (GP63 or leishmanolysin), one of *Leishmania* virulence factors
- During differentiation process, expression of certain genes occur that preadapt the parasites to enter and survive in macrophage phagolysosome.



PCR-RFPL profile of Gp63



• 2 restriction enzymes using *SalI* (A) and *BglI* (B)

Leishmania (subgenus Viannia)

l = *L. (V.) lainsoni*; g = *L. (V.) guyanensis*; b = *L. (V.) braziliensis*;
p = *L. (V.) peruviana* M= Molecular Marker

Real-Time PCR



- *Compare the traditional leishmania diagnostic methods (culture and microscopy) versus RT-PCR in cutaneous leishmaniasis

Sensitivities

■ RT-PCR	95% (95% CI; 90-98%)
■ Culture	78% (95% CI; 99-86%, P=0.001)
■ Microscopy	73% (95% CI; 63-81% P <0.0005)

Molecular Diagnosis for Parasitic Diseases

Diagnostic technique	Identification of the parasite				
	PCR	Culture	Direct search	Serology	
Simple	+	-	++	+/-	
Sensitive	+++	++	-	++	Accuracy in diagnosis
Specific	+++	+++	+++	++	
Fast	+++	-	++	+++	Early diagnosis
Cost efficient	+/-	-	+++	+/-	applicability

Leishmaniasis, Chagas disease, onchocercosis, toxoplasmosis, asymptomatic malaria

*Dept. of Tropical Medicine, Instituto Oswaldo Cruz, Fundação Oswaldo Cruz
Ministry of Health, Rio de Janeiro, Brasil*



Comparison of 3 PCR assays for Diagnosis of Cutaneous Leishmaniasis

-kDNA gene (the highest sensitivity)	98.7% (77/78)
- ITS1 of SSu r RNA gene	91.0% (71/78)
- Miniexon gene	53.8% (42/78)
- Microscopy	83.3% (65/78)



Species identification

- Multilocus enzyme electrophoresis (Isoenzyme) analysis is the gold standard

Disadvantages

- requires parasite isolation and cultivation
- Time-consuming
- Require technical expertise



Species identification

- When parasite isolation and cultivation could not be available, a simplified and specific method for species diagnosis and identification of *Leishmania* is needed
- Using PCR, a minute amount of DNA is enough to amplify well-known marker genes by this technique.



Species identification

■ PCR-RFLP

–ITS1 of SSU rRNA gene; restriction enzyme

Hae III

–Mini-exon gene; restriction enzyme, *Eae I*

■ DNA sequencing of PCR products

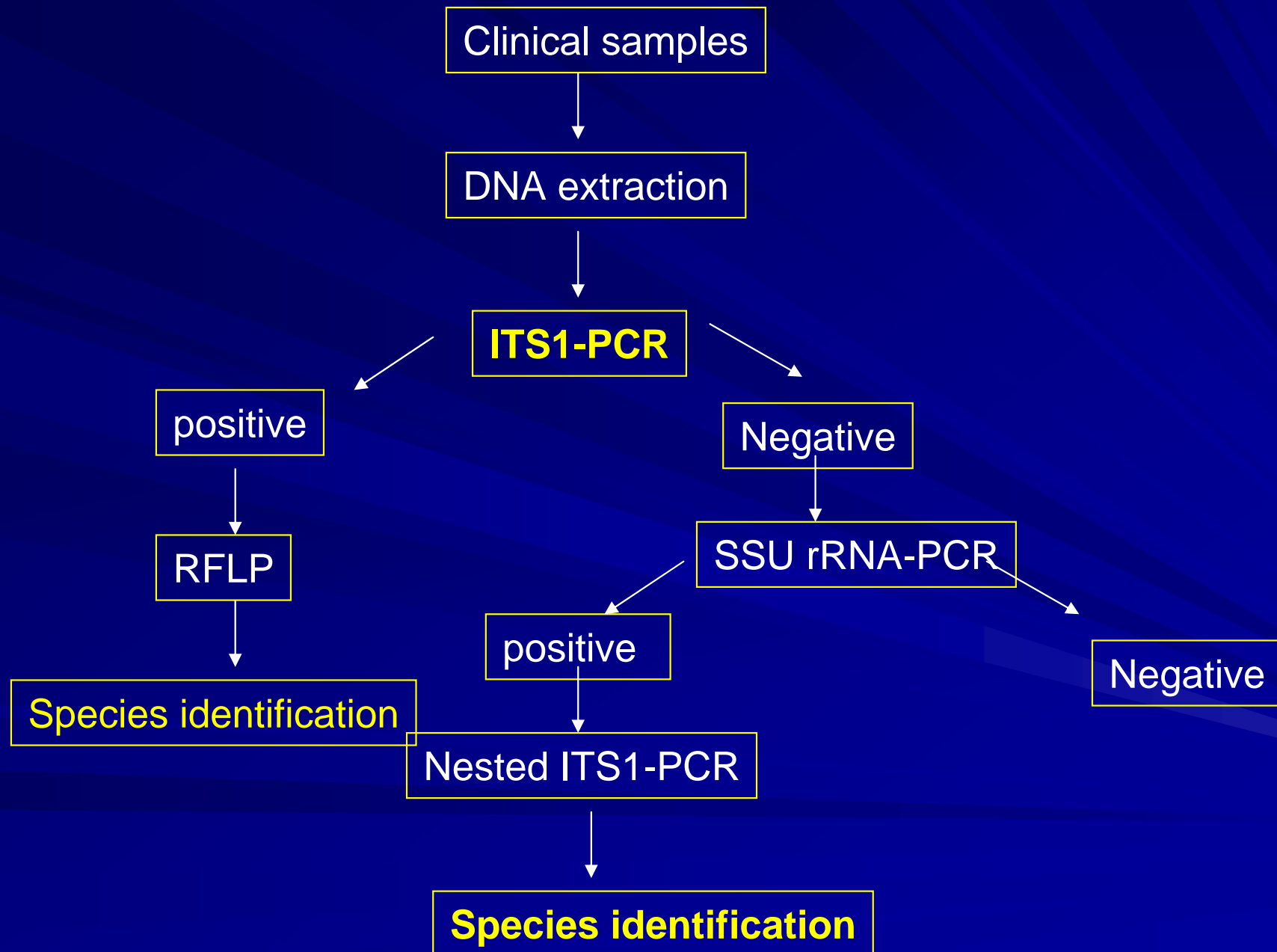
Rotureau B, et al., J Clin Microbiol. 2006;44:459–67.

Marfurt J. et al., Diagn Microbiol Infect Dis. 2003;46:115-24.



Species identification (ITS1*)

by Schonian G et al., Diag Microbiol Inf Dis 2003; 47: 349-358



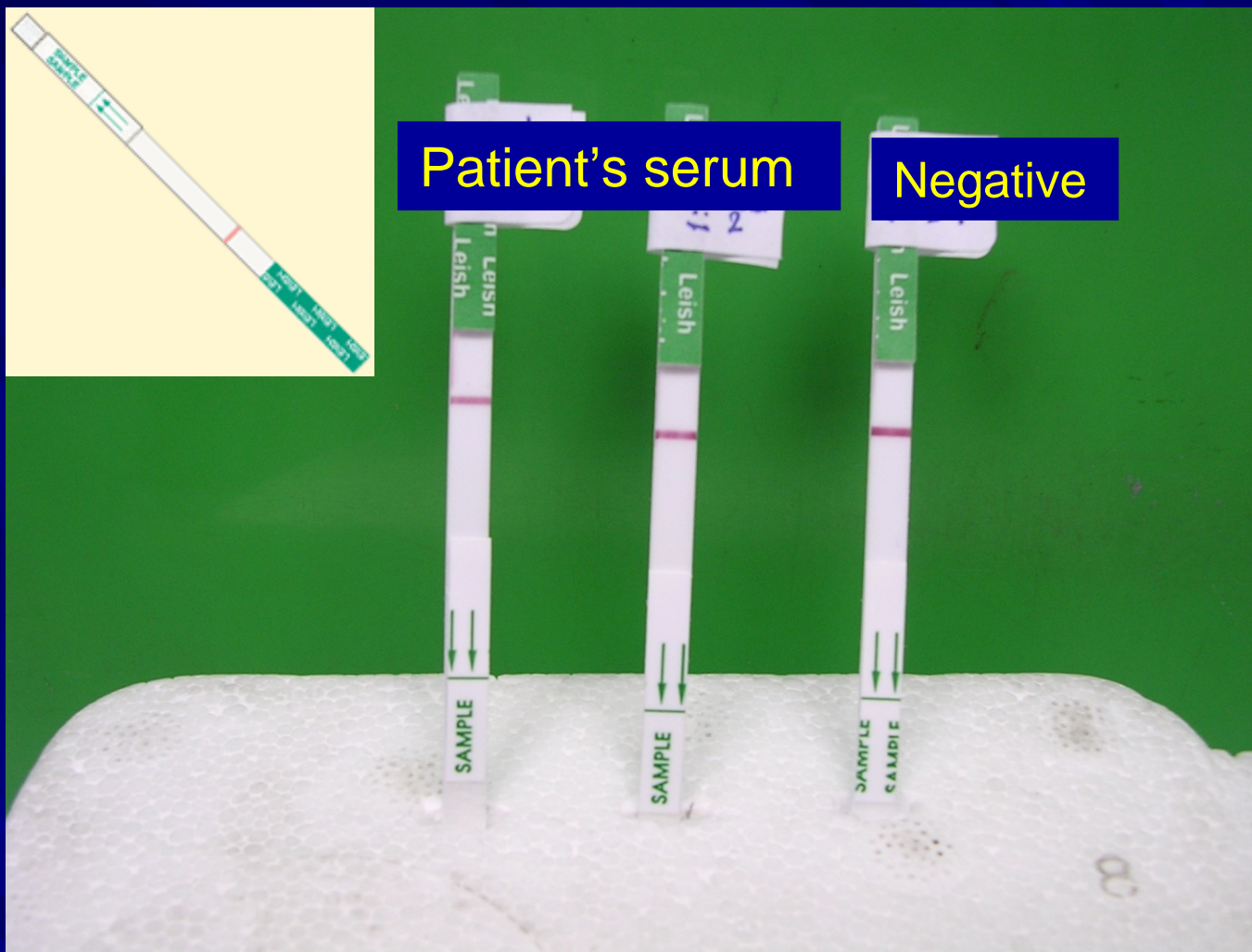
An autochthonous Visceral leishmaniasis, Phangnga province



- No VL cases were previously reported in this province.
- Bone marrow smear : positive amastigotes
- DAT : positive, titer 1:200
- rk39 : **negative**



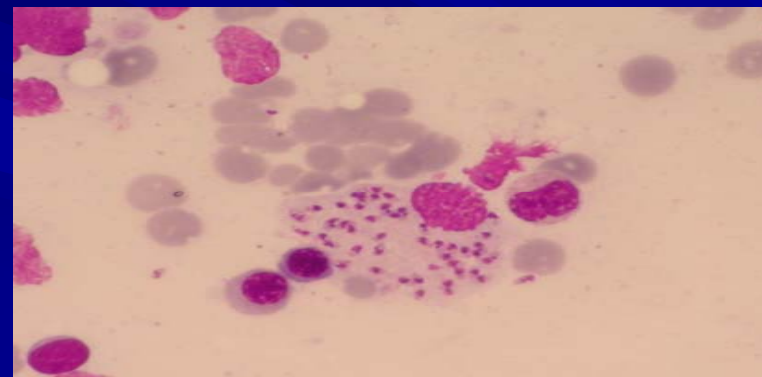
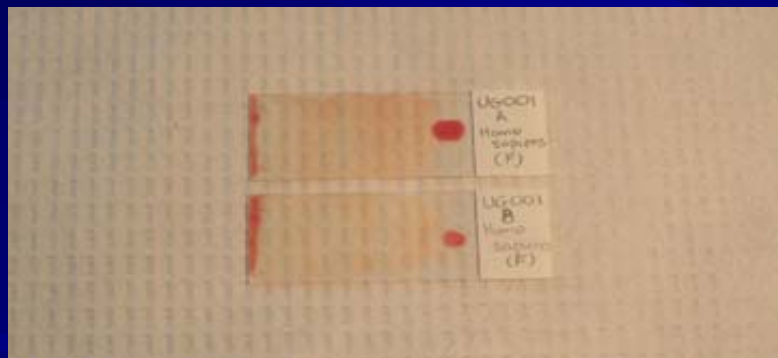
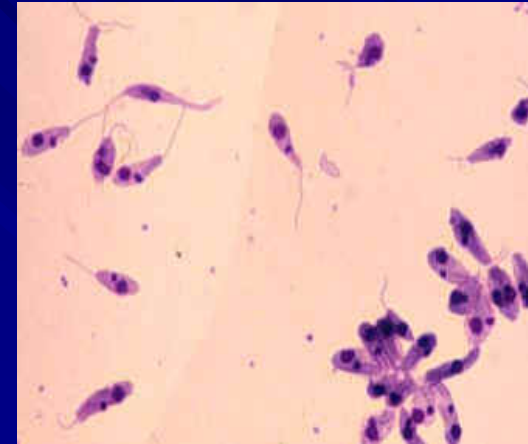
rk39 Strip Test





Samples

- Promastigotes from cultivation
- Giemsa-Stained bone marrow smears on glass slides

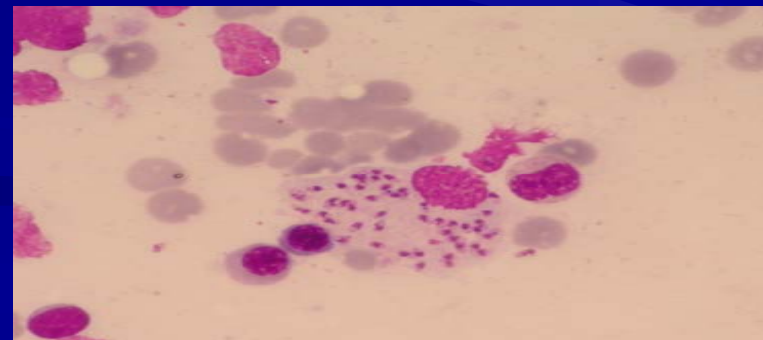
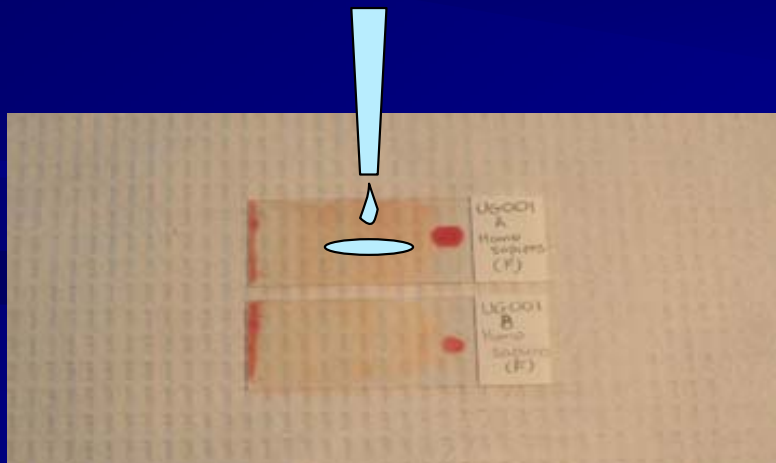




MATERIALS AND METHODS

1. Samples

- Giemsa-Stained bone marrow smears on glass slides



MATERIALS AND METHODS

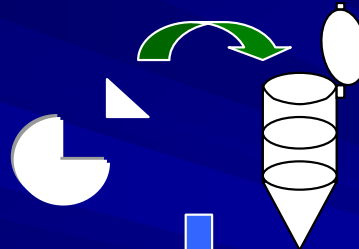


2. DNA Extraction



14 μ l of specimen was applied on to the FTA disk

Air dried



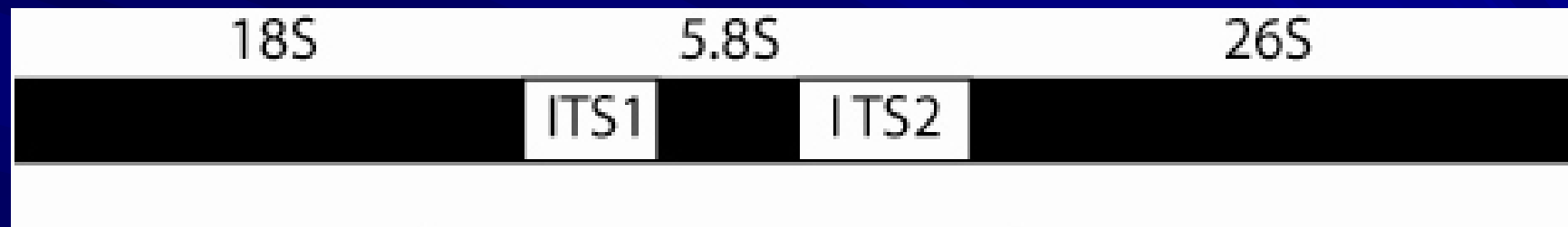
Washed twice with FTA purification reagent and washed twice again with TE buffer

Dried at 80°C for 2 hr.

MATERIALS AND METHODS



3. Amplification of SSU-rRNA / ITS1 region



SSU rRNA gene, primers

S4 : 5'-GAT CAA GCT GCA GGT TCA CC-3'

S12 : 5'-GGT TGA TTC CGT CAA CGG AC-3'

ITS1, primers

LITSR : 5'-CTG GAT CAT TTT CCG ATG -3'

L5.8S : 5'-TGA TAC CAC TTA G TCG CAC TT-3'

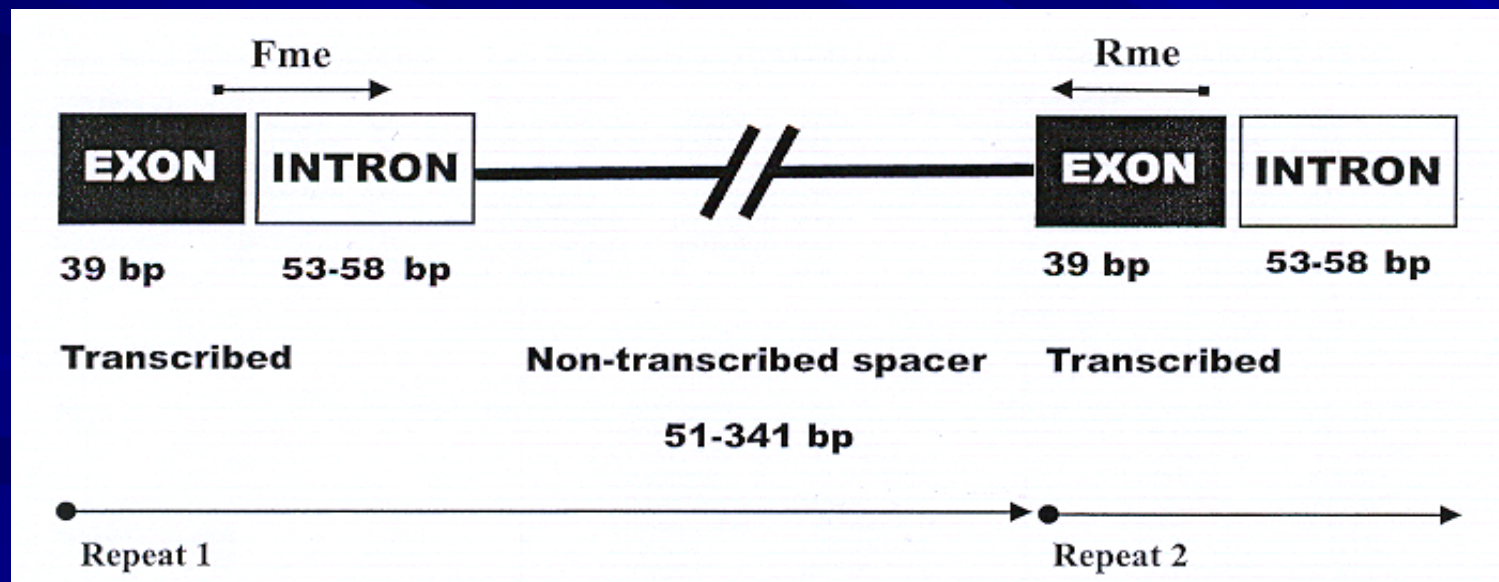


MATERIALS AND METHODS

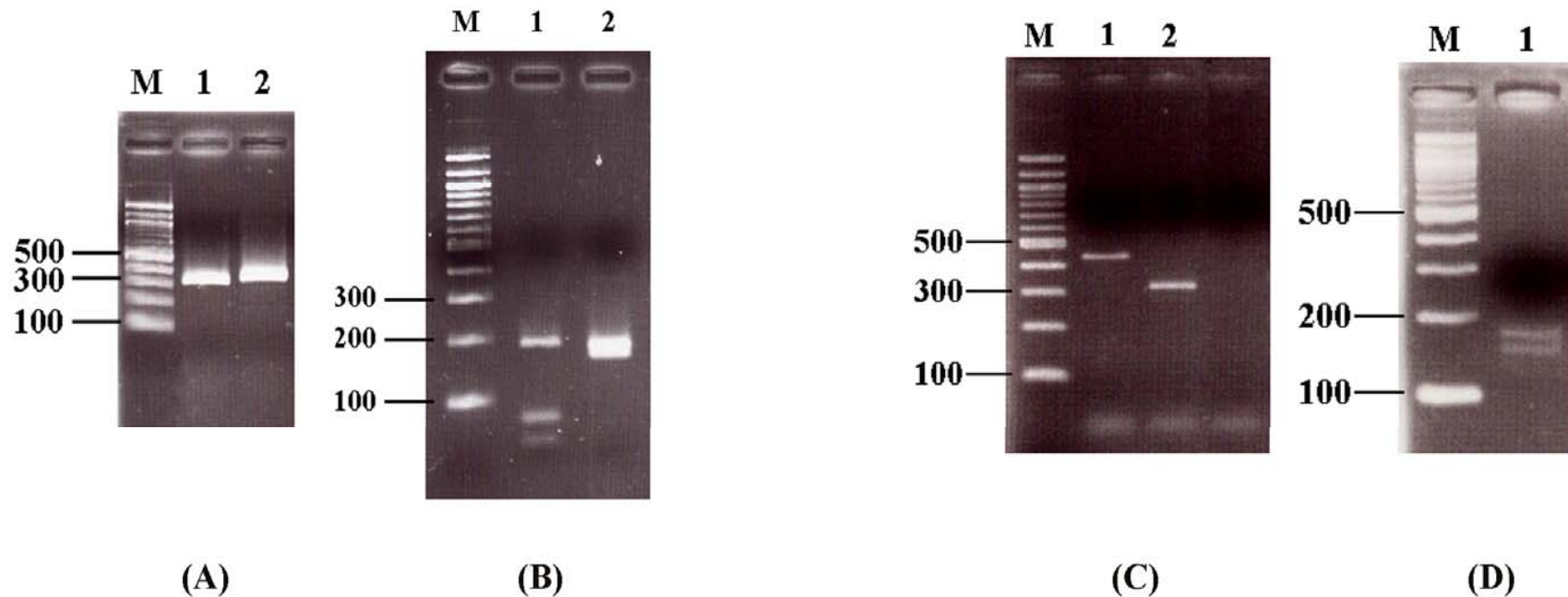
■ Mini-exon gene amplification, primers

Fme : 5'-TAT TGG TAT GCG AAA CTT CCG -3'

Rme : 5'-ACA GAA ACT GAT ACT TAT ATA GCG -3'



Marfurt J, et al., J Clin Microbiol. 2003;41:3147-53.



1 = positive control *L. donovani*;

2 = *Leishmania* unknown

A) PCR amplification of ITS1 region of SSU-RNA gene

B) RFLP patterns of ITS1 region of SSU-RNA gene digest with *HaellI*

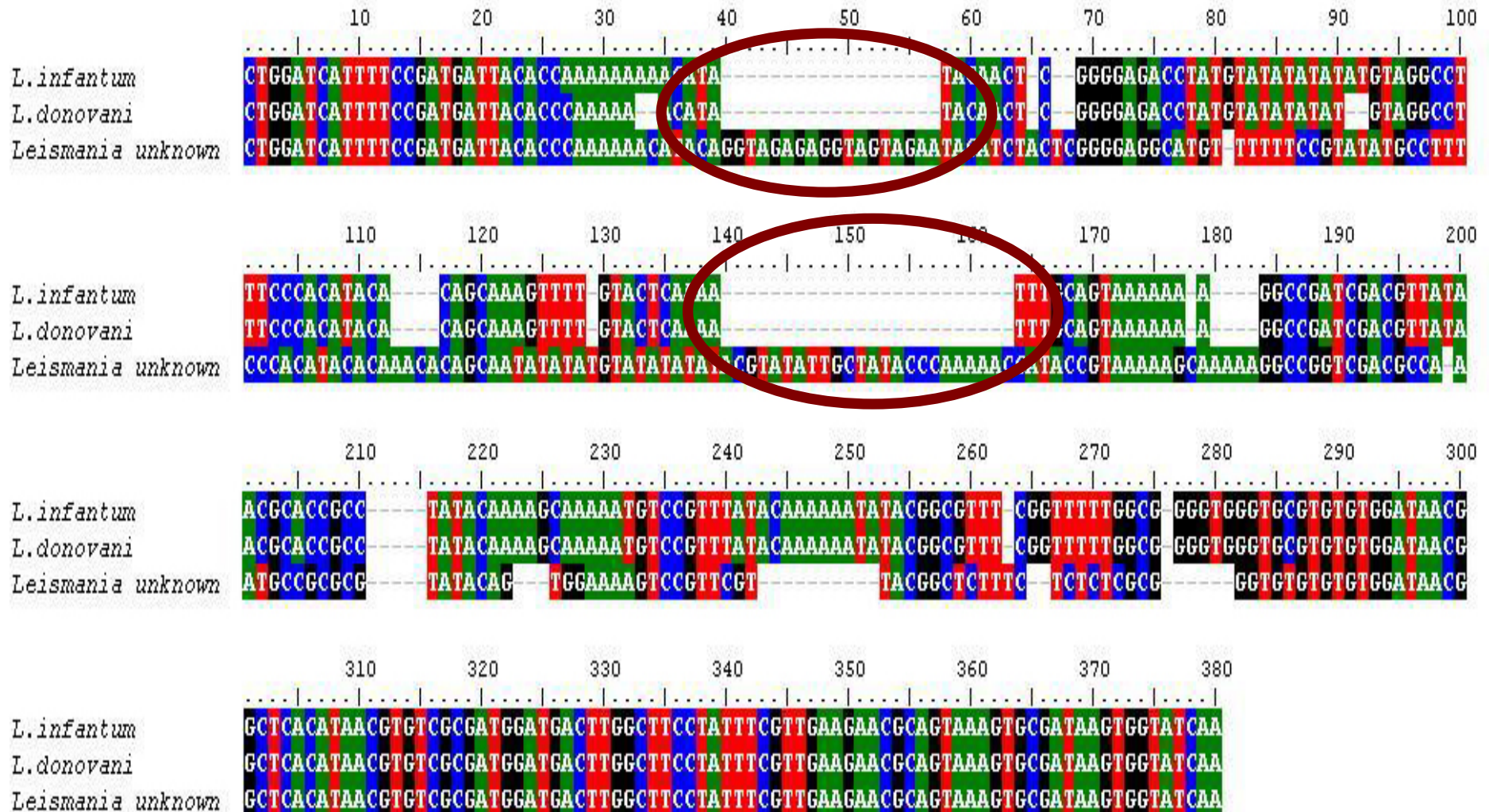
C) PCR amplification of mini-exon region of SSU-RNA gene

D) RFLP patterns of mini-exon gene of *Leishmania* unknown with *EaeI*.

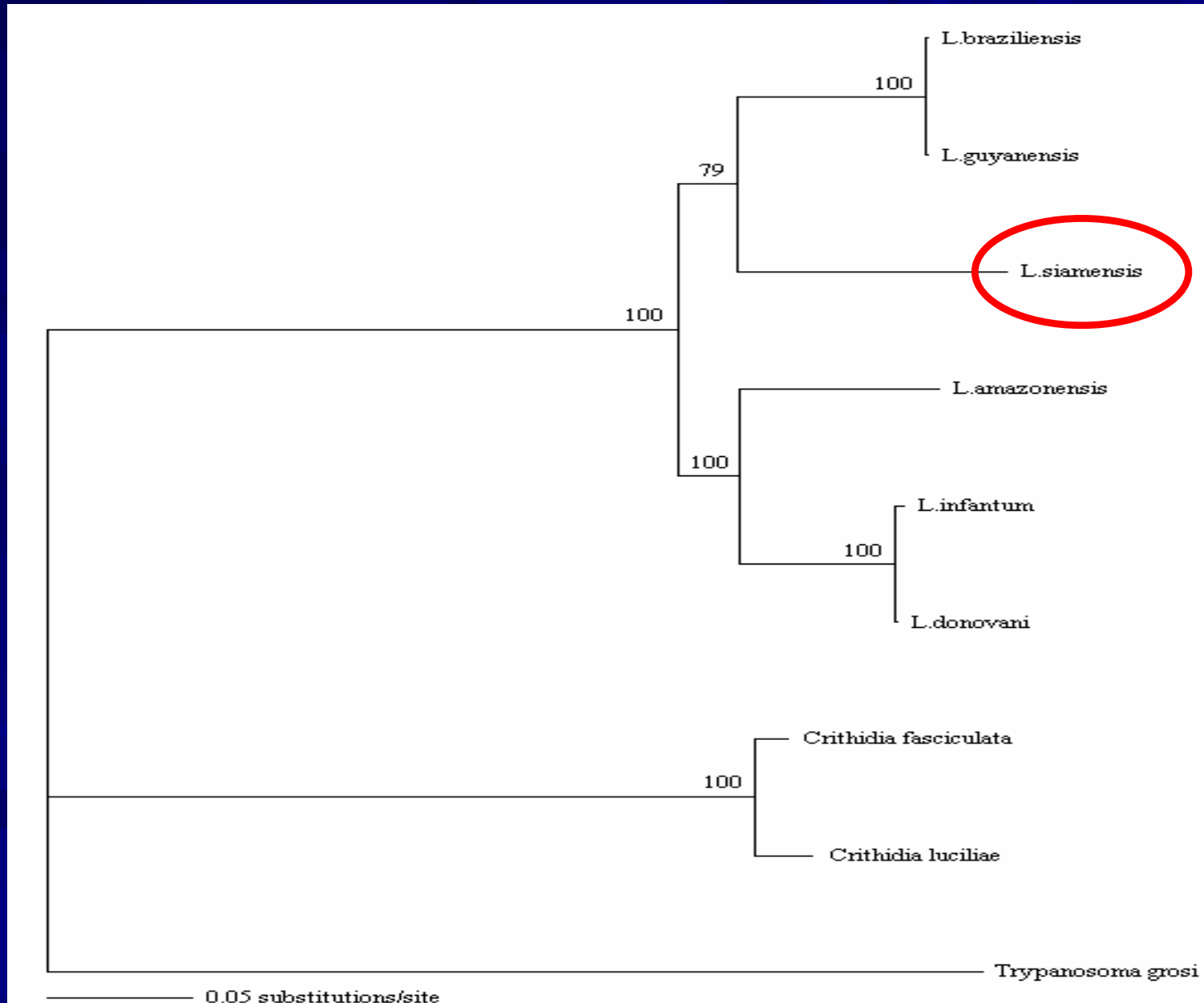
```
L. infantum CTGGATCATTTTCCGATGATTACACCAAAAAAAAAACATA-----TAC
L. donovani CTGGATCATTTTCCGATGATTACACCCAAAAA--ACATA-----TAC
L. amazonensis CTGGATCATTTTCCGATGATTACACCAAAAAACATATA-----CAA
L. braziliensis CTGGATCATTTTCCGATGATTACACCAAAAAA-----CA-----TAC
L. guyanensis CTGGATCATTTTCCGATGATTACACCAAAAAA-----CA-----TAC
Leishmania. unknown CTGGATCATTTTCCGATGATTACACCAAAAAACATACAGGTTAGAGAGGTAGTAGAATAC
***** * * * * *
L. infantum AACT-C--GGGGAGACCTATGTATATATATATGTAGGCCTTCCACATACA----CAGC
L. donovani AACT-C--GGGGAGACCTATGTATATATAT--GTAGGCCTTCCACATACA----CAGC
L. amazonensis AACT-CTCGGGGAGACCTAT-TCTTTTCGATAGGCAGCCTTCCACACATACA----CAGC
L. braziliensis AACT-C-CGGGGAGGCTTGTGTTTTTCTAGCAAGC--CTTTCCACAGATACG----CAAT
L. guyanensis AACT-C-CGGGGAGGCTTGTGTTTTTCTAGCAAGC--CTTTCCACAGATACG----CAAT
Leishmania. unknown ATCTACTCGGGGAGGCATGT-TTTTTCCGTATATGCCTTCCACATASACAAACACAGC
* * * * * * * * * * * * * * * *
L. infantum AAAGTTTT-GTACTCAAAA-----TTTTGCAGTAAAAAA-A-
L. donovani AAAGTTTT-GTACTCAAAA-----TTTTGCAGTAAAAAA-A-
L. amazonensis AAAGTTTTGTTACTCAAAA-----CAACATTTGCAGTAAACAA-AA
L. braziliensis ACAATCTATATAATATATAT-----ATATAGACACAACATACAGTAGAAAA-A-
L. guyanensis ACAATCTATATAATATATATGT-----ATATAGACACAACATACAGTAGAAAA-A-
Leishmania. unknown AATATATATGTATATATATACGTATATTGCTATACCCAAAACCATACCGTAAAAAGCAA
* * * * * * * * * * * * * * * *
L. infantum ---GGCCGATCGACGTTATAACGCACCGCC-----TATACAAAAGCAAAAATGTCCGTTT
L. donovani ---GGCCGATCGACGTTATAACGCACCGCC-----TATACAAAAGCAAAAATGTCCGTTT
L. amazonensis AATGGCCGATCGACGTTATAGCGCACAACCGCGTATATACAAAAGCAGAGAAAAATGCC
L. braziliensis ---GGCCGATCGACGTTA--ACATATCGCG-----TATACAA---CAAAAAGTTCCGTT
L. guyanensis ---GGCCGATCGACGTTA--ACATATCGCG-----TATACAA---CAAAAAGTTCCGTT
Leishmania. unknown AAAGGCCGTCGACGCCA-AATGCCGCGCG-----TATACAG---TGGAAAAGTCCGTT
***** * * * * * * * * * *
L. infantum ATACAAAAAATATACGGCGTTT-CGGTTTTTGGCG-GGGTGGGTGCGTGTGTGGATAACG
L. donovani ATACAAAAAATATACGGCGTTT-CGGTTTTTGGCG-GGGTGGGTGCGTGTGTGGATAACG
L. amazonensis GTTTCAA-----TACGGCGTTTTCCGTTTTTGGGGCGGGGGGTGCGTGTGTGGATAACG
L. braziliensis -----TACGGCTTTT-----TTTTGGCG-----GCGTGCCTGGATAACG
L. guyanensis -----TACGGCTTTT-----TTTTGGCG-----GCGTGCCTGGATAACG
Leishmania unknown GT-----TACGGCTCTTTT--TCTCTCGCG-----GGTGTGTGTGTGGATAACG
***** * * * * * * * * * *
L. infantum GCTCACATAACGTGTGCGGATGGATGACTTGGCTTCCCTATTTTCGTTGAAGAACGCAGTAA
L. donovani GCTCACATAACGTGTGCGGATGGATGACTTGGCTTCCCTATTTTCGTTGAAGAACGCAGTAA
L. amazonensis GCTCACATAACGTGTGCGGATGGATGACTTGGCTTCCCTATTTTCGTTGAAGAACGCAGTAA
L. braziliensis GCTCACATAACGTGTGCGGATGGATGACTTGGCTTCCCTATTTTCGTTGAAGAACGCAGTAA
L. guyanensis GCTCACATAACGTGTGCGGATGGATGACTTGGCTTCCCTATTTTCGTTGAAGAACGCAGTAA
Leishmania unknown GCTCACATAACGTGTGCGGATGGATGACTTGGCTTCCCTATTTTCGTTGAAGAACGCAGTAA
*****
L. infantum AGTGCGATAAGTGGTATCAA
L. donovani AGTGCGATAAGTGGTATCAA
L. amazonensis AGTGCGATAAGTGGTATCAA
L. braziliensis AGTGCGATAAGTGGTATCAA
L. guyanensis AGTGCGATAAGTGGTATCAA
Leishmania unknown AGTGCGATAAGTGGTATCAA
*****
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Neighbor-joining based on ITS1 region sequence analysis containing 6 species.

Alignment of ITS1 region



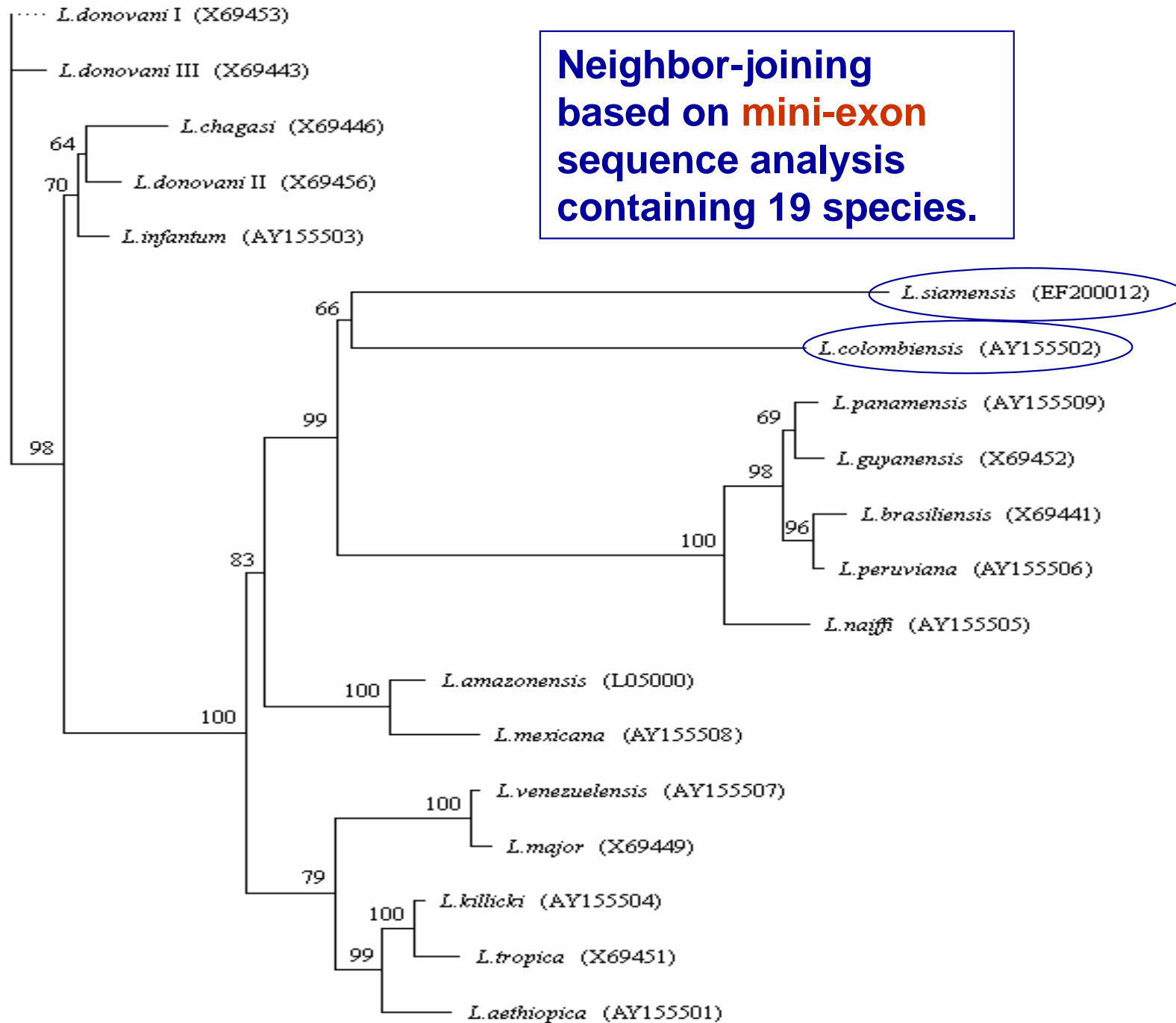
Phylogenetic tree, ITS1



Percent identity of ITS1 sequences comparison between unknown *Leishmania*, and other species

Pair-wise comparison	% Identity of ITS1
<i>L. braziliensis</i> vs. <i>Leishmania</i> unknown	83
<i>L. guyanensis</i> vs. <i>Leishmania</i> unknown	84
<i>L. amazonensis</i> vs. <i>Leishmania</i> unknown	74
<i>L. infantum</i> vs. <i>Leishmania</i> unknown	77
<i>L. donovani</i> vs. <i>Leishmania</i> unknown	78
<i>L. braziliensis</i> vs. <i>L. guyanensis</i>	100
<i>L. braziliensis</i> vs. <i>L. amazonensis</i>	79
<i>L. braziliensis</i> vs. <i>L. infantum</i>	82
<i>L. braziliensis</i> vs. <i>L. donovani</i>	81
<i>L. guyanensis</i> vs. <i>L. amazonensis</i>	79
<i>L. guyanensis</i> vs. <i>L. infantum</i>	81
<i>L. guyanensis</i> vs. <i>L. donovani</i>	81
<i>L. amazonensis</i> vs. <i>L. infantum</i>	90
<i>L. amazonensis</i> vs. <i>L. donovani</i>	90
<i>L. infantum</i> vs. <i>L. donovani</i>	99

Neighbor-joining
based on mini-exon
sequence analysis
containing 19 species.



0.05 substitutions/site

Alignment of the 320 basepairs from the sequenced PCR products of
mini-exon gene of *L. siamensis* (EF200012) and *L. colombiensis* (AY155502).

```

L.colombiensis      TTTTGGGAAGCGCGCAAGCGCTACATTTTTTTTTTTTG---TCATGTGCGGG-----TGCG
Leishmania Unknown TTTTGG-AGCGCGCCAGCGCTCTCTTTTTTTTTTTTG-GTGC-TGCGCGTG-----TGCA
*****  *****  *****  **********  *  **  ***  *      ***

L.colombiensis      CGCCGCCCTCCTCCACTGGGGGGGCTTCCCACACGCGCTGTGGTCCCTTTTCCCCTTG
Leishmania Unknown CGCCCACGCGCCGGCCCCCTCCACGGCGGGTGC GGCGCCCGGCTGCTGCTCGACGGC---
****  *  *  **  *      *      ***      ***  *      **  *  *

L.colombiensis      ----CTGCTGGGAGGGGGG-GGGGCGCGCCGCTGCGGGTGGTTTCCTCCTGTGCGGCGTCC
Leishmania Unknown -----CGCCGCT-----GCCCCTCACGCCCGGCG-CC
                *  *                        *      ***** **

L.colombiensis      CCTCGGGCCTGCCCGGCAT-CCTTGCTGGGGGCTCTGCCGGCGCGATTCCGGCACCCCAT
Leishmania Unknown CGTCCGGCGGG-CACGCGCCGCC-CAGCCG-----GCCCCG-----GCAGTCCGCCAT
*  *****  *  *  **      *  *  *  *      ***  **      *  *  *  ****

L.colombiensis      GCCCAGTGGCCGGGACCCCCCCTCTGGTCCCCCCCCCATGATCCGGCGCACGCCTGACAC
Leishmania Unknown GACGCGCG-CCTCGGCGCAGCCAC-----C
*  *  *  *  **  *  *  *  **

*

L.colombiensis      GCGGTTAGTGCA-TGCCAAATGAGCCACCC
Leishmania Unknown ACCGCCACCCCCCAAACG-GCGCGCCGGCT
                *  *  *  *      *  ***  *

```

54% consensus identity.

***A novel species of *Leishmania*, *Leishmania siamensis*, sp. nov.**



Conclusion

- DAT and rK39 dipstick show similar diagnostic performance for serological studies in the field.
- The high sensitivity and ease of performance make both DAT and rK39 dipstick very suitable for surveillance surveys. However it is essential to evaluate any rapid diagnostic tests in the region in which it will be used.
- PCR-based protocols have increased the speed and sensitivities of species-specific leishmaniasis diagnosis.
- A novel species of *Leishmania* caused visceral leishmaniasis was identified in 1 autochthonous Thai patient, southern Thailand.
- Efforts should be made PCR platforms more user-friendly and cost-effective for epidemiological studies.

**Department of Parasitology,
Phramongkutklao College of Medicine**



Saovanee leelayoova
Mathirut Mungthin
Suradej Siripattanapipong
Tawee Naaglor

**Department of Biotechnology, Faculty of Science and
Institute of Science and Technology for Research and
Development, Mahidol University**

Jeerapun Worapong

**Field Epidemiology Training Program - Thailand, Bureau
of Epidemiology, Ministry of Public Health**

Teerayuth Sukmee