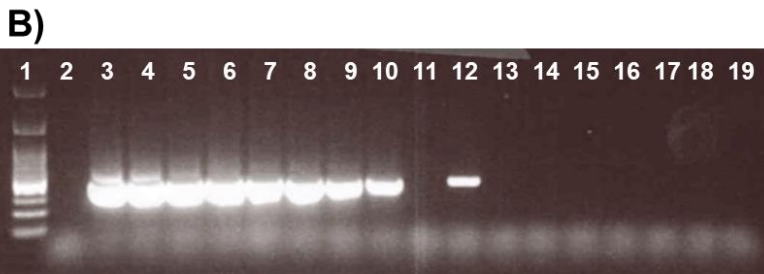
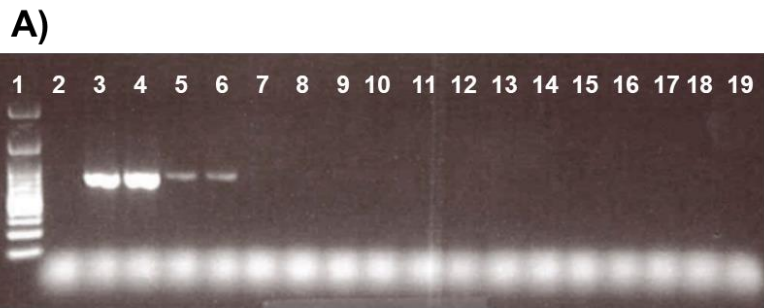
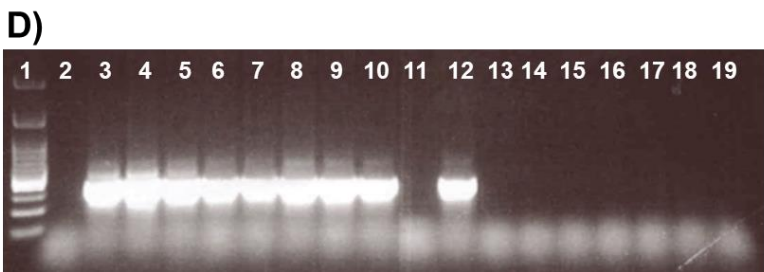
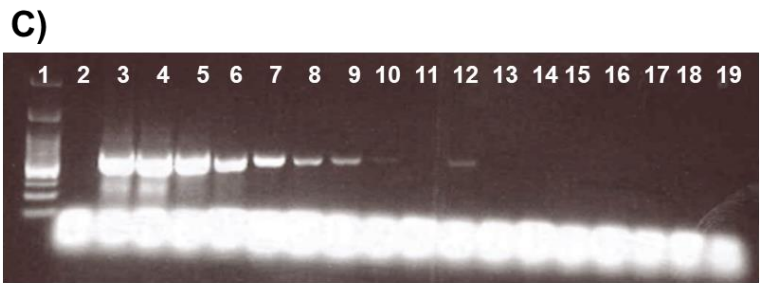


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## SUPPLEMENTAL MATERIAL

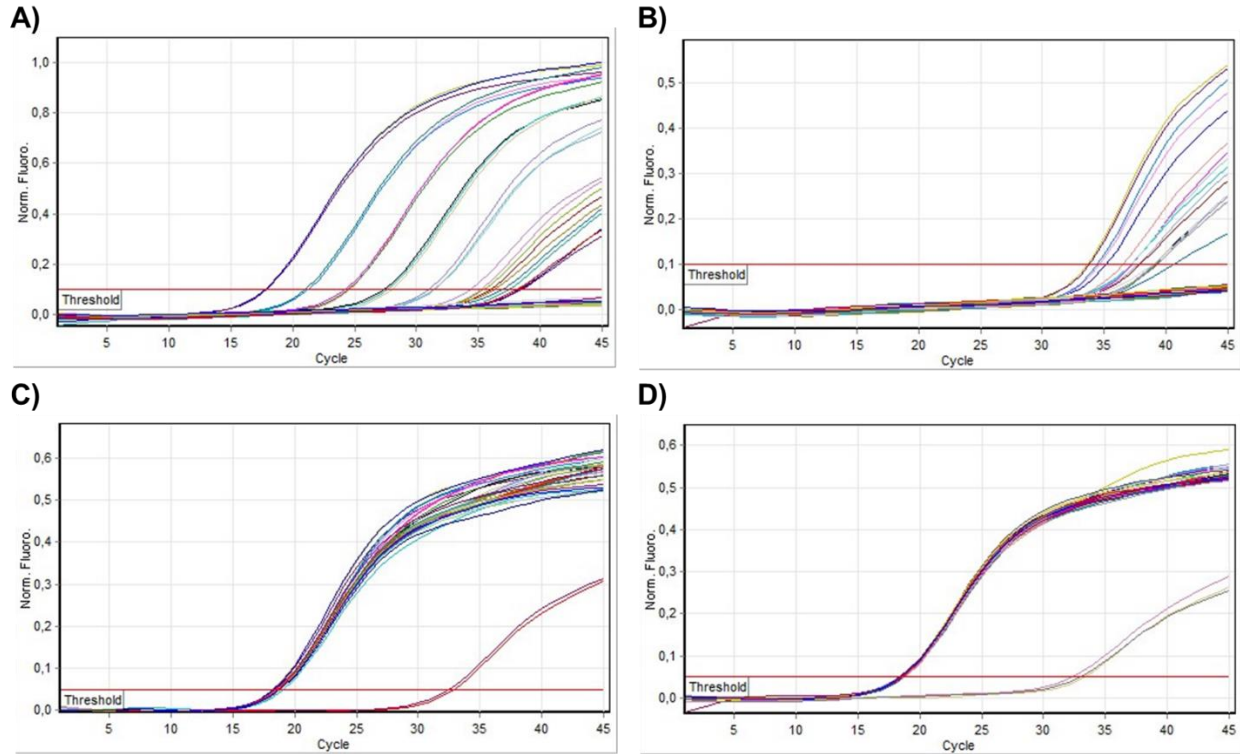


2



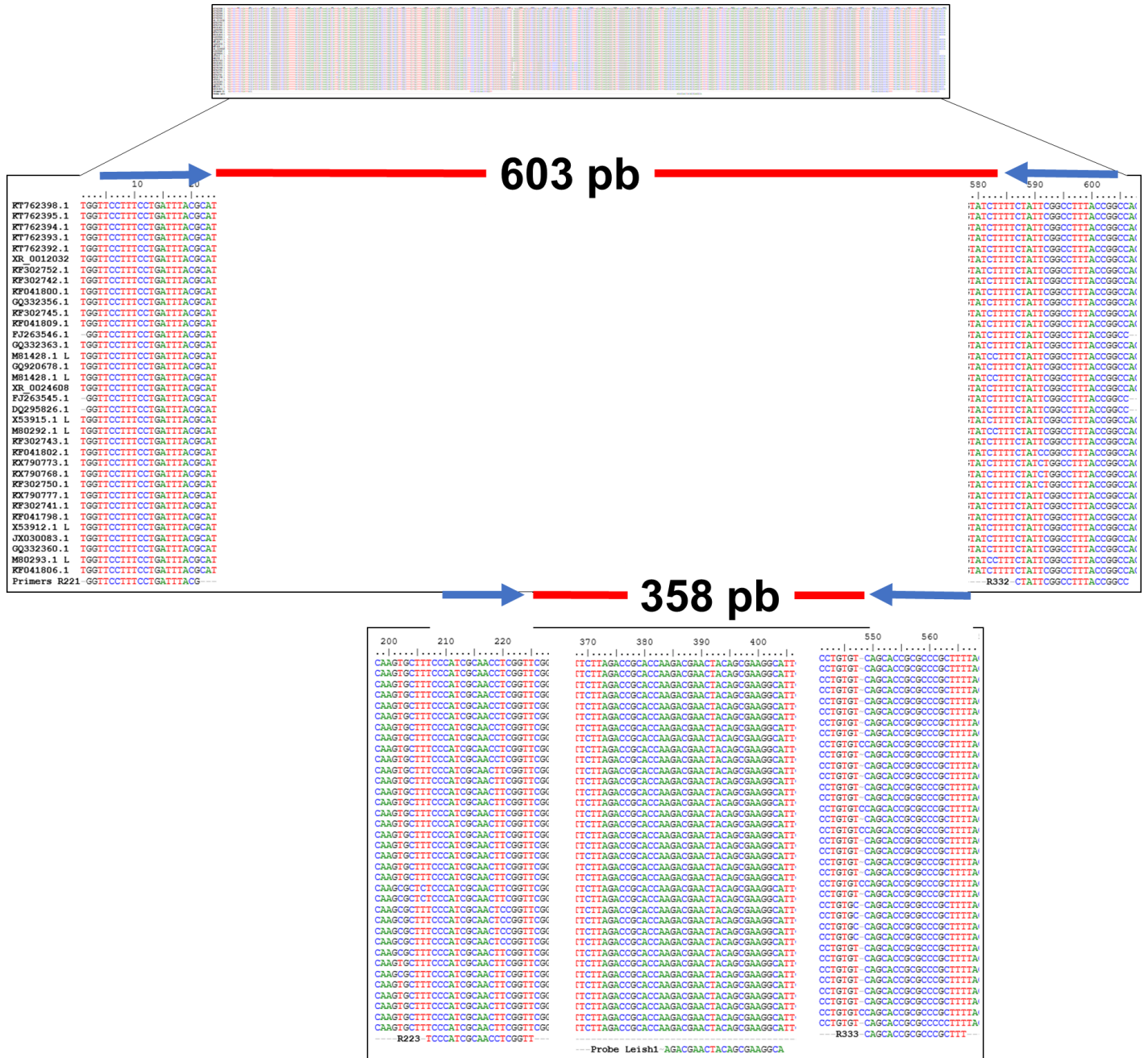
3

4 **Figure S1.** Analytical sensitivity of first and second PCR reaction of Ln-PCR (A & B) and  
5 LeishGelPCR (C & D) using spiked samples from  $5 \times 10^6$  to 0.05 parasites/reaction. Line 1, 100 bp  
6 ladder; line 2, not template control; line 3,  $5 \times 10^6$ ; line 4,  $5 \times 10^5$ ; line 5,  $5 \times 10^4$ ; line 6,  $5 \times 10^3$ ; line 7,  
7 500; line 8, 50; line 9, 5; line 10, 0.5; line 11, 0.05 parasites/reaction; line 12, positive control, the  
8 size of amplified product is 600 bp in the first reaction, and 330 bp in the second reaction.



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**Figure S2.** Analytical sensitivity of Leish-qPCR using spiked samples: A) Blood samples with DNA *Leishmania* adjusted at  $1 \times 10^6$  to 0.001 parasites equivalents/reaction tested in triplicate. B) Blood samples with *Leishmania* DNA adjusted at 1 to 0.01 parasites equivalents/reaction tested in quintuple. C & D) Detection of the internal control.



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**Figure S3.** Alignment of different *Leishmania* species. Sequences were downloaded from Gene bank. Sequences of R333 and R332 are in reverse complementary. Superior part is the product can be obtained in the first reaction of Ln-PCR and LeisGelPCR. Lower part is the product of the second reaction of nested PCRs. Also, it is the product of the Leish-qPCR. *L. chagasi*: KT762398.1; KT762395.1; KT762394.1; KT762393.1; KT762392.1. *L. infantum*: XR\_001203206.1; KF302752.1. *L. donovani*: KF302742.1; KF041800.1; GQ332356.1. *L. tropica*: KF302745.1; KF041809.1; FJ263546.1; GQ332363.1. *L. aethiopia*: M81428.1; GQ920678.1; M81428.1. *L. major*: XR\_002460808.1; FJ263545.1; DQ295826.1; X53915.1. *L. brasiliensis*: M80292.1. *L. guyanensis*: KF302743.1. *L. equatorensis*: KF041802.1; KX790773.1. *L. colombiensis*: KX790768.1; KF302750.1. *L. enriettii*: KX790777.1; KF041798.1. *L. mexicana*: KF302741.1. *L. amazonensis*: X53912.1; JX030083.1. *L. mexicana*: GQ332360.1; M80293.1; KF041806.1.

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**Table S1.** Comparison of the limit of detection of Ln-PCR and LeishGelPCR

| N parasites per reaction | Ln-PCR                                    |   | LeishGelPCR                               |   |
|--------------------------|---|---|---|---|
|                          | 1st reaction                              | 2nd reaction                              | 1st reaction                              | 2nd reaction                              |
|                          | N Pos reactions per total repetitions (%) | N Pos reactions per total repetitions (%) | N Pos reactions per total repetitions (%) | N Pos reactions per total repetitions (%) |
| <b>5</b>                 | 0/10 (0)                                  | 10/10 (100)                               | 8/10 (80)                                 | 10/10 (100)                               |
| <b>0.5</b>               | 0/10 (0)                                  | 8/10 (80)                                 | 4/10 (40)                                 | 10/10 (100)                               |
| <b>0.05</b>              | 0/10 (0)                                  | 4/10 (40)                                 | 0/10 (0)                                  | 1/10 (10)                                 |

Pos, positive.

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**Table S2.** Limit of detection (LOD) of Leish-qPCR

| Parasites per reaction | Average of C <sub>T</sub> | Positive reactions per total repetitions (%) | Expected positive reactions | Likelihood (%) |
|------------------------|---------------------------|--|-----------------------------|----------------|
| <b>1</b>               | 34.66                     | 8/8 (100)                                    | 8                           | 100            |
| <b>0.1</b>             | 37.23                     | 8/8 (100)                                    | 7                           | 91.6           |
| <b>0.01</b>            | 37.90                     | 5/8 (62.5)                                   | 6                           | 76.5           |
| <b>0.001</b>           | 39.26                     | 2/8 (25)                                     | 2                           | 20.9           |

36 At 95% of probit likelihood, the LOD of Leish-qPCR in terms of C<sub>T</sub> value was 37.084 (CI 95%: 34.33 – 37.74), so in  
37 terms of parasites/reaction, the LOD was 0.2 (CI 95%: 0.1 – 1.1).  
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39 **Table S3.** Comparison of Ln-PCR, LeishGelPCR and Leish-qPCR using clinical samples from  
40 leishmaniasis patients.

| Sample |        | Ln-PCR       |       |                    |                    | LeishGelPCR  |       |                    |                    | Leish-qPCR |
|--------|--------|--------------|-------|--------------------|--------------------|--------------|-------|--------------------|--------------------|------------|
| Code   | Matrix | 1st reaction |       | 2nd reaction       |                    | 1st reaction |       | 2nd reaction       |                    |            |
|        |        | 10 μL        | 20 μL | 10 μL <sup>a</sup> | 20 μL <sup>a</sup> | 10 μL        | 20 μL | 10 μL <sup>a</sup> | 20 μL <sup>a</sup> |            |
| S01    | FTB    | P            | P     | P                  | P                  | P            | P     | P                  | P                  | P          |
| S02    | LNA    | P            | P     | P                  | P                  | P            | P     | P                  | P                  | P          |
| S03    | BM     | P            | P     | P                  | P                  | P            | P     | P                  | P                  | P          |
| S04    | WB     | P            | P     | P                  | P                  | P            | P     | P                  | P                  | P          |
| S05    | BM     | P            | P     | P                  | P                  | P            | P     | P                  | P                  | P          |
| S06    | BM     | N            | N     | P                  | P                  | P            | P     | P                  | P                  | P          |
| S07    | FTB    | N            | N     | P                  | P                  | P            | P     | P                  | P                  | P          |
| S08    | WB     | N            | N     | P                  | P                  | P            | P     | P                  | P                  | P          |
| S09    | FTB    | N            | N     | P                  | P                  | N            | N     | N                  | P                  | P          |
| S10    | FTB    | N            | N     | N                  | P                  | N            | N     | P                  | P                  | P          |

41 BM, bone marrow; FTB, fresh tissue biopsies; LNA, lymph node aspirate; WB, whole blood; P, positive; N, negative.

42 <sup>a</sup>The 2nd reaction was carried out with 10 μL of the 1:40 dilution of the amplified product of the 1st reaction.

43 **Table S4.** Comparison of Ln-PCR, LeishGelPCR and Leish-qPCR using DNA samples from  
 44 uninfected individuals and other pathogens

| Clinical status                   | Matrix of sample | N samples | N positive reactions by Ln-PCR | N positive reactions by LeishGel PCR | N positive reactions by Leish-qPCR |
|-----------------------------------|------------------|-----------|--------------------------------|--------------------------------------|------------------------------------|
| Uninfected                        |                  |           |                                |                                      |                                    |
| CL suspicion                      | FTB              | 2         | 0                              | 0                                    | 0                                  |
| VL suspicion                      | BM               | 4         | 0                              | 0                                    | 0                                  |
|                                   | WB               | 1         | 0                              | 0                                    | 0                                  |
|                                   | LNA              | 1         | 0                              | 0                                    | 0                                  |
| Other parasite infections         |                  |           |                                |                                      |                                    |
| <i>Cryptosporidium sp</i>         | Stool            | 2         | 0                              | 0                                    | 0                                  |
| <i>Giardia lamblia</i>            | Stool            | 2         | 0                              | 0                                    | 0                                  |
| <i>Blastocystis hominis</i>       | Stool            | 2         | 0                              | 0                                    | 0                                  |
| <i>Dientamoeba fragilis</i>       | Stool            | 2         | 0                              | 0                                    | 0                                  |
| <i>Plasmodium falciparum</i>      | WB               | 2         | 0                              | 0                                    | 0                                  |
| <i>Toxoplasma gondii</i>          | CSF              | 1         | 0                              | 0                                    | 0                                  |
| <i>Trypanosoma cruzi</i>          | WB               | 2         | 0                              | 0                                    | 0                                  |
| <i>Mycobacterium tuberculosis</i> | Culture          | 2         | 0                              | 0                                    | 0                                  |

45 BM, bone marrow; CSF, cerebrospinal fluid; FTB, fresh tissue biopsies; LNA, lymph node aspirate; WB, whole blood.