Development of a novel loop-mediated isothermal amplification assay for rapid detection of *Mycobacterium leprae* in clinical samples

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Abstract

Background: Sensitive and definitive diagnostic tests are required for timely treatment of leprosy and to control its transmission. **Aim:** In the present study, we report the development of loop-mediated isothermal amplification assay using six primers targeting the *RLEP* gene sequence uniquely present in *Mycobacterium leprae*.

Methods: Tissue punch samples (*n* = 50) and slit aspirates (*n* = 50) from confirmed cases of leprosy (*M. leprae* positive by quantitative polymerase chain reaction), reporting at the Department of Dermatology, Safdarjung Hospital, New Delhi, were analyzed using newly developed closed tube loop-mediated isothermal amplification assay. The sensitivity and specificity; positive predictive value, negative predictive value and accuracy were calculated using MedCalc statistical software.

Results: The loop-mediated isothermal amplification assay specifically amplified *M. leprae* genomic DNA with an analytical sensitivity of 100 fg. About 47 Out of the 50 quantitative polymerase chain reactions confirmed *M. leprae* positive tissue samples, 47 were positive by loop-mediated isothermal amplification assay (sensitivity 94%; 95% confidence interval 83.5%–98.8%) while only 31/50 were positive by histopathology (sensitivity 62%; 95% confidence interval 47.2%–75.4%). Using slit aspirate samples of these 50 patients, 42 were positive by both quantitative polymerase chain reaction and loop-mediated isothermal amplification assay (sensitivity 84%; 95% confidence interval 31.8%–60.7%) were positive by both polymerase chain reaction and loop-mediated isothermal amplification assay (sensitivity 84%; 95% confidence interval 31.8%–60.7%) were positive by microscopy.

Limitations: In the present study, the leprosy patient cohort was not uniform, as it comprised a lower number of paucibacillary cases (22%) compared to multibacillary (78%) cases.

Conclusion: Loop-mediated isothermal amplification assay established here provides a rapid and accurate diagnostic test for leprosy in terms of sensitivity and specificity. The assay is simple to perform in comparison with other molecular techniques (polymerase chain reaction/quantitative polymerase chain reaction) and has potential for field applicability.

Key words: Diagnosis, histopathology/microscopy, leprosy, loop-mediated isothermal amplification, quantitative polymerase chain reaction

Introduction

Leprosy caused by *Mycobacterium leprae*, an acid-fast bacterium, has been known since the biblical times with reports of cases dating over 3000 years ago.¹ The disease is endemic in tropical countries with new cases being mainly

reported from India (58%), Brazil (16%) and Indonesia (9%).² According to the World Health Organization, the global prevalence at the end of 2016 was 171,948 with a registered prevalence rate of 0.23/10,000 individuals.³ In India, the leprosy prevalence had decreased markedly since

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the introduction of multidrug therapy, from a prevalence rate of 57.8/10,000 in 1983 to 0.84/10,000 in 2006; however, the rate of new case detection has increased.⁴ According to the National Leprosy Elimination Program, Government of India (2018), about 88,000 people in India are suffering from leprosy (prevalence rate = 0.66/10000) with 135,485 new cases reported in 2017 (annual new case detection = 10.17/10000).

The spread of leprosy is usually caused by frequent close contact for a longer duration between an untreated patient and a person genetically susceptible to developing the disease. Patients often experience numbness, skin lesions, muscle weakness, enlarged nerves and joint pain. It has been observed that leprosy is often misdiagnosed with other similar skin diseases of the infiltrated plaque, papules or nodules including cutaneous leishmaniasis, macular postkala-azar dermal leishmaniasis, cutaneous tuberculosis, sarcoidosis, lymphoma and syphilis.⁵ Accurate, specific and sensitive methodologies are needed to provide a definite diagnosis. Presently, microscopic observation of Ziehl-Neelsen stained slides6,7 and histopathological examination of cutaneous lesion biopsy^{8,9} are considered as "gold standard" for the diagnosis of leprosy but are less sensitive, time-consuming and dependent on technical expertise. The molecular methodologies, although not routinely used, are sensitive and specific; however, these need sophisticated and expensive equipment that may not be available in laboratories with limited resources. There have been numerous studies based on molecular techniques such as polymerase chain reaction or nested polymerase chain reaction to detect M. leprae in tissue samples¹⁰⁻¹³ and in slit aspirate samples^{10,14,15} with variable sensitivity and specificity. The highly conserved repetitive sequence RLEP has been the preferred target due to its high copy number (n = 37),^{10,14} although other targets such as rpoT, 16S rRNA and Sod A have also been exploited for detection of *M. leprae* by polymerase chain reaction/ quantitative polymerase chain reaction.^{16,17} Several cost-effective isothermal amplification-based techniques have emerged to substitute expensive molecular methods.¹⁸ Amongst these, loop-mediated isothermal amplification is simple, rapid, specific and sensitive and only a heating block or water bath capable to maintain a constant temperature (60 °C to 65 °C) is required.¹⁹ The technique has many advantages such as the reaction proceeds isothermally,¹⁹ crude DNA extracts can be used directly without purification,^{19,20} and the products can be detected visually using multiple parameters including turbidity, fluorescence and color. For naked-eye detection, intercalating dye such as SYBR Green I, calcein or malachite green is added to the amplified products.^{21,22} The present study is focused to develop leprosy specific loop-mediated isothermal amplification assay targeting the RLEP gene to detect M. *leprae* in clinical samples.

Methods

Parasite DNA samples

To examine the specificity of the designed primers, *M. leprae* genomic DNA (NR-19350), obtained from BEI Resources, VA, USA, was tested along with extracted DNA from various coendemicdisease-causingbacteria/parasitessuchas*Mycobacterium tuberculosis, Acinetobacter baumannii, Klebsiella pneumoniae, Leishmania donovani, Leishmania major, Leishmania tropica, Plasmodium vivax and Plasmodium falciparum* used as negative controls. Once the novel loop-mediated isothermal amplification assay for *M. leprae* detection was established, it was applied to the clinical samples of leprosy patients, while samples from other skin diseases were used as controls to determine the diagnostic specificity of the assay.

Clinical samples

The study was conducted after obtaining ethical clearance under the guidelines of the International Ethics Committee of Safdarjung Hospital, New Delhi and informed consent of the participants. At pretreatment stage, 3 mm punch tissue sample from skin lesions (n = 50) and slit aspirates taken from standard sites i.e., bilaterally at the eyebrows, earlobes and additionally selected site that observed skin lesions (n = 50) in 200µL NET buffer (150 mmol/L NaCl, 15 mmol/L Tris-HCl [pH 8.3] and 1 mmol/L EDTA) were collected from leprosy patients presenting characteristic clinical manifestations, reporting at the dermatology department of our hospital from June 2017 till June 2018. All cases were confirmed by quantitative polymerase chain reaction using tissue samples. Tissue samples from other skin diseases including post-kala-azar dermal leishmaniasis, vitiligo, sporotrichosis and pityriasis lichenoides chronica, lichen sclerosis, pityriasis rosea (n = 25) and normal skin samples (n = 15) were used as negative controls. Patients positive for HIV, hepatitis B and C, tuberculosis or any other systemic ailments were excluded. Pregnant or lactating women were excluded from the study. For DNA isolation, QIAamp DNA mini kit was used following manufacturer's instructions. The isolated tissue DNA were eluted in 50µL ($\approx 170 \text{ ng/µL}$) and slit aspirate DNA in 20µL ($\approx 3.5 \text{ ng/µL}$) of nuclease-free water and stored at -30°C till further use.

Quantitative real-time polymerase chain reaction

The quantitative polymerase chain reaction was performed using primers described in an earlier study [Table 1 and Figure 1].¹¹ Briefly, *M. leprae* specific real-time polymerase chain reaction was set up in an ABI Prism 7500 sequence detection system (Applied Biosystems, USA) using *RLEP* based forward and reverse primers. The quantitative polymerase chain reaction was performed as per the manufacturer's instructions in a 10 μ L reaction mixture consisting of 1X SYBR Green Fast polymerase chain reaction Master mix (Applied Biosystems, USA), 5 pmol forward primer, 5 pmol reverse primers and 1 μ L volume of DNA from the sample. The results were obtained in accordance with the first fluorescent signal detection cycle threshold and the sample was considered positive when it

Table 1: Primer sequences for SYBR green I-based qPCR and loop-mediated isothermal amplification assay			
Name of primer	Primer sequence (5'-3')	Reference	
qPCR primer			
RLEP-forward	TGCATGTCATGGCCTTGAGG	11	
RLEP-reverse	CACCGATACCAGCGGCAGAA		
LAMP primer			
RLEP-F3	TTGTTGGTGGGTGGCTGA	This study	
RLEP-B3	CGGCGCTAACAACTATCCTC		
RLEP-FIP (F1c-F2)	TTACGTGCGCCGCGCTAATCCTGCTTTCGATGAGGCTTCG		
RLEP-BIP (B1c-B2)	GGTGGATGCTGCTTGGTCTACATGCATCGATATCGCCTTCAG		
RLEP-FLP	CACTGCGGCAAAGCACA		
RLEP-BLP	TGTTGATGATGCCAGGGGC		

FIP: forward inner primer, BIP: backward inner primer, F3: outer forward primer, B3: outer backward primer, FLP: forward loop primer, BLP: backward loop primer, qPCR: quantitative polymerase chain reaction, LAMP: loop-mediated isothermal amplification

1 TGANCANCCC TTACAGGTCG AGCTCGAGGG TTTCTTGAAG CATCGAGGTC CTACCACGTG 61 GTTGCCAACT TGGCCTGAGT GGCGTCAAAA ATCGTGCGGT TCCCGGGCTC TGCTGTCTTG 121 TGTCTGTGGC GGTGAGTTGT TCGGTGGTGT TCATAGGTGG TGGGGTGAAAT GGCTGTTTTT 181 GCGTTTTATG ACTGGCCGAT ATGTTCGGTA GTCGTGGGGG GCAGCCCGGA ATCCTGTTGA 241 CGTGTTTGC TGTGTGCGG GGTTTTTGTT GGTGGGGGGC TGACTGCCTG CTTTCGATGA 241 CGTGTTTGC TGTGTGCGC AGTGGACACG ATTAGCGGGG CGCACGTAG CATGCCGGTG F2 FLP 301 GGCTTCGTGT CCTTGGCCGC AGTGGACACG ATTAGCGGGG CGCACGTAG GCATGCGGTG 51 GTGGATGCTG CTTGGTCAC ATGTTGATGA TGCCAGGGC TGGGCACCTG GGCTGCGGG 51 GTGGATGCTG CTTGGTCAC ATGTTGATGA TGCCAGGGC TGGGCACCTG GGCTGGCGG 52 FLP 421 AAGGCGATAT CGATGCAGGC GTGAGTGGTG GGATAGTTGT TAGCGCCGCG GGGTAGGGGC 52 B3 481 GTTTTAGTG <u>GCATGCATG GCCTGACG</u> GTCGGCGTGG TCAATGTGGC CGCACCTGAA 781 CAGGCACGTC CCCGTGCACG GTATAACTAT TCGCACCTGA TGTTATCCCT TGCACCATT 601 <u>CTGCCCGCGGG TACCGGTGC</u> GGCGGCTTGT TGACCGGCCC TCAGCCAGCA AGCAGGCATG 721 CACATACGGC AACCTTCTAG GGCGGCTTGT TGACCGGCCC TCAGCCAGCA AGCAGGCAGG 721 CACATACGGC AACCTTCTAG CGCAGATCA CCACCACAC CCCACCACCA CACACACA 781 CCACCCAAAC CAAACCAGCA AAAAATAACC ACCAAATGAC CATCACGACG ACACCTTGAA 841 GTCTAAAACGG ACAAGTTACT GGTTACTGGC CGGGCGGGC TTTACCCGAT AGCAGGGAC 901 GGCTGTCAAC GCGATACTT AGCCCCCCAT GGCAGATGA ATCATAGTCA GCTCGGGAA 901 GGCTGTCAAC GCGATACTT AGCCCCCCAT GGCAGATGAA ATCATAGTCA GCTCGGGTAA 1021 GAACGATCGA GTCATTCCTA GACCCCAAAC ACCACACGAT GTCGATCAGG CACTGAATGC 1081 TCGGCAGGAC CAAAATAACC TCCTTTGCC TTAAGATTC	>X17153.1 <i>M. leprae</i> repetitive element, RLEP3				
61 GTTGCCAACT TGGCCTGAGT GGCGTCAAAA ATCGTGCGGT TCCCGGGCTC TGCTGTCTG 121 TGTCTGTGGC GGTGAGTTGT TCGGTTGGTT TCATAGGTGG TGGGTGAAAT GGCTGTTTT 181 GCGTTTTATG ACTGGCCGAT ATGTTCGGTA GTCGTGGGGG GCAGCCCGGA ATCCTGTTGA 241 CGTGTTTTGC TGTGTTGCGG GGTTTTTGT GGTGGGGGGG TGACTGCCTG CTTTCGATGA 241 CGTGTTTGC TGTGTTGCCGC AGTGGACACG ATTAGCGGCGG CGCACCTAG CATGCGGTG F2 FLP FLP F1c 301 GGCTTCGTGT CTTGGTCTAC ATGTTGATGA TGCCAGGGGC TGGGCACCTG GGCTGTGCTG B1c BLP 421 AAGGCGATGCT CTGGTCTAC ATGTTGATGA TGCCAGGGGC TGGGCACCTG GGCTGGCGG B2 B2 B3 481 GTTTTAGT <u>G GCATGCCAGG</u> GTGAGTGTGA GGATAGTTGT TAGCGCCGGG GGTAGGGGC B2 B3 481 GTTTTAGT <u>G GCATGCCAGG</u> GTGAGTGTGA GGATAGTTGT TAGCGCCGCG GGGTAGGGGC B2 B3 481 GTTTTAGT <u>G CCATGCCAGG</u> GTGATAACTAT TCGCACCTGA TGTTATCCCT TGCACCATTT 601 <u>CTGCCGCGGG TACCGCTGC</u> GGCGGCTTGT TGACCGGCCC TCAGCCAGCA AGCAGGCATG RLEP-F 541 CAGGCACGTC CCCGTGCAGG GTATAACTAT TCGCACCTGA TGTTATCCCT TGCACCATTT 601 <u>CTGCCGCGGG</u> GAGCAGTATC GTGTTAGTGA ACAGTGCAC CACCACAGCA AGCAGGCATG RLEP-R 661 CCGCCGGGGT CAGCAGTATC GTGTTAGTGA ACAGTGCAC CACCACAGCA AGCAGGCAG 721 CACATACGGC AACCTTCTAG CGCAGATCAA CCACCCACAC CCCACCAGCC CACCACAACA 781 CCACCCAAAC CAAACCAGCA AAAAATAACC ACCAAATGAC CATCACGACG ACACCTTGAA 841 GTCTAAACGG ACAGTTACT GGTTACTGGC CGGCGCGGC TTTACCCGAT AGCAGGCGGC 901 GGCTGTCAAC GCGATACTTC ACGCCCCCAT GGCCAAGGAG ATCATAGTCA GCTCGGGGTAA 1021 GAACGATCGA GTCTATCCGA CTTCGCCCT GGCAAGAGA ATCATAGTCA GCTCGGGTAA 1021 GAACGATCGA GTCTATCCAG CCCCCAAAC ACCACACGAT GTCGATCAGG CACTGGATAG 1081 TCGGCAGGAC CAAAAGTAGC TCCTCTTCGC TTAAGGAT TC	1 TGAACAACCC TTACAGGTCG AGCTCGAGGG TTTCTTGAAG CATCGAGGTC CTACCACGTG				
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1081 TCGGCAGCAC CARARGTAGC TCCTCTTCGC TTARGAATTC	1021 GAACGATCGA GTCTATCCTA GACCCCAAAC ACCACGAT GTCGATCACG CACTGAATGO				

Figure 1: *RLEP* gene (Accession no X17153.1) sequence showing six primers (outer forward primer, outer backward primer, forward inner primer, backward inner primer, forward loop primer and backward loop primer) for loop-mediated isothermal amplification assay; Forward and reverse primers for quantitative polymerase chain reaction

showed cycle threshold smaller than 34 (negative cutoff value of >34). The cycling parameters included 50 °C for 2 min, 95 °C for 10 min and 40 cycles of 95 °C for 15 s and 61 °C for 60 s. The analytical sensitivity of SYBR Green I based *RLEP* specific quantitative polymerase chain reaction

assay was determined using serially diluted (tenfold; 1 ng/ μ L to 1 fg/ μ L) *M. leprae* genomic DNA.

Loop-mediated isothermal amplification assay

Loop-mediated isothermal amplification assay was developed targeting repetitive and specific RLEP gene (accession no X17153.1) sequence uniquely present in M. leprae. The six primers; forward inner primer, backward inner primer, outer forward primer, outer backward primer, forward loop primer and backward loop primer were designed using Primer Explorer V4 software (http://primerexplorer. jp/e/) [Table 1 and Figure 1]. Loop-mediated isothermal amplification reaction was performed as described earlier²² in 25µL reaction mixture containing 40 pmol each of forward inner primer and backward inner primer primers, 5 pmol each of outer forward primer and outer backward primer primers, 20 pmol each of the forward loop primer and backward loop primer, 1.4 mM of each deoxynucleoside triphosphate, 0.8 M betaine, 20 mM Tris-HCl, pH 8.8, 10 mM KCl, 10 mM (NH₄)₂SO₄, 8 mM MgSO₄, 0.1% TritonX-100, 8U Bst 2.0 (New England Biolabs, Massachusetts, USA) and 1-5 µL sample DNA at different time points (30 min to 120 min) at 65 °C in a heating block. For naked-eye detection, closed tube technology was used where SYBR Green I (Molecular Probes, Eugene, OR, USA) was added inside the tube cap at the beginning and a quick spin was given after completion of the assay to allow mixing of SYBR Green I with the amplified product. Green color in tube indicated positive while negative remained orange. Once the assay was established it was applied to clinical samples to determine diagnostic sensitivity of the assay. The assay was performed at 65 °C for varied time points ranging from 30 min to 120 min for different clinical samples. The reaction time for loop-mediated isothermal amplification assay was optimized as 60 min for tissue samples and 90 min for slit aspirate samples.

Statistical analysis

Samples were confirmed as *M. leprae* positive using quantitative polymerase chain reaction as a "gold standard" and classified as true positive, true negative, false positive and false negative for loop-mediated isothermal amplification

assay evaluation. The clinical sensitivity, specificity, 95% confidence interval and accuracy were calculated for loopmediated isothermal amplification using MedCalc statistical software. The positive predictive value was calculated as (number of true positives)/(number of true positives + number of false positives) ×100 and the negative predictive value was calculated as (number of true negatives)/(number of true negatives) ×100. The accuracy was calculated as (number of true positives + number of true positives + number of true positives + number of true negatives) ×100. The accuracy was calculated as (number of true positives + number of true positives + number of true positives + number of true negatives) ×100.

Results

The average age of patients (n = 50) was 36 years (14–72 years) with men: women ratio is 2.8:1 (37 men and 13 women patients). Amongst the leprosy patients, on the basis of the number of lesions present all over the body, 39 patients were characterized as multibacillary presenting more than five lesions, while 11 as paucibacillary with less than five lesions, at the time of sample collection.

Analytical sensitivity and specificity of quantitative polymerase chain reaction

The quantitative polymerase chain reaction was performed as a confirmatory assay to determine *M. leprae* positive samples. The limit of detection for *RLEP* based quantitative polymerase chain reaction assay was 10 fg that is equivalent to approximately three *M. leprae* organisms¹³ and the standard curve showed reproducibility with a negative correlation between cycle threshold and DNA concentration of *M. leprae* [Figure 2a]. To evaluate the specificity of the quantitative polymerase chain reaction assay, control parasite DNA samples along with *M. leprae* genomic DNA were subjected to the quantitative polymerase chain reaction. The assay specifically detected *M. leprae* DNA. The cycle threshold value for all control samples was higher than the negative control (negative cutoff value of >34) indicating 100% specificity of the assay [Figure 2b].

Analytical sensitivity and specificity of loop-mediated isothermal amplification assay

The loop-mediated isothermal amplification assay was performed using serially diluted *M. leprae* genomic DNA samples (tenfold; Ing/ μ L to 1fg/ μ L). The analytical sensitivity of loop-mediated isothermal amplification in 60 min was 100 fg, equivalent to 30 organisms [Figure 3a]. To evaluate the specificity of the loopmediated isothermal amplification assay, parasite DNA controls along with *M. leprae* genomic DNA were subjected to the loopmediated isothermal amplification reaction. The assay detected *M. leprae* DNA as indicated by green color while all the control samples remained orange after completion of the reaction, suggesting that no amplification occurred (specificity- 100%; 95% confidence interval 92.9%–100%) [Figure 3b].

Application of loop-mediated isothermal amplification assay in clinical samples

The loop-mediated isothermal amplification assay was applied to clinical samples and its efficacy was compared with the routinely used diagnostic tests (histopathology/microscopy) for M. leprae detection. Out of the 50 M. leprae positive tissue samples as confirmed by quantitative polymerase chain reaction, 47 tissue samples produced green color after 60 min incubation in loop-mediated isothermal amplification assay (sensitivity 94%; 95% confidence interval 83.5%-98.8%), a positive predictive value of 100% and negative predictive value of 93% (95% confidence interval 81.7%-97.6%) with an accuracy of 96.6% (95% confidence interval 90.5%-99.3%) with the assay being negative in all 25 other disease control samples, giving it a specificity of 100% (95% confidence interval 86.28% to 100.00%), while only 31 were positive for M. leprae in histopathology reports (sensitivity 62%; 95%) confidence interval 47.2%–75.%) [Table 2 and Figure 4a]. Amongst these tissue samples, 37 out of 39 multibacillary samples (sensitivity 94.9%; 95% confidence interval 82.7%-99.4%) and ten out of 11 paucibacillary samples (sensitivity 90.9%; 95% confidence interval 58.7%-99.8%) were positive by loop-mediated isothermal amplification assay whereas only 26 multibacillary samples (sensitivity 66.6%; 95% confidence interval 51%-79.4%) and five paucibacillary samples (sensitivity 45.5%; 95% confidence interval 21.3%-72%) were found positive in histopathology reports [Table 2].



Figure 2a: Limit of detection for *M. leprae* at 10 fold dilution $(1 \text{ ng}/\mu\text{L}-1 \text{ fg}/\mu\text{L})$ by quantitative polymerase chain reaction

In the present study, we attempted to establish loop-mediated isothermal amplification assay for leprosy using slit aspirates



Figure 2b:Specificity of quantitative polymerase chain reaction primers for detection of M. leprae was tested using DNA from other organisms.



Figure 3: (a) Visual detection of amplified loop-mediated isothermal amplification products using SYBR green I. Sensitivity for *M. leprae* (1 ng/ μ L-1 fg/ μ L), (b) Visual detection of amplified loop-mediated isothermal amplification products using SYBR green I

of patients, as this sample collection procedure is far less invasive with advantages of ease of collection, storage and transportation. The slit aspirate samples of these patients showed 42/50 (sensitivity 84%; 95% confidence interval 70.9%-92.8%) positives after 90 min incubation by loop-mediated isothermal amplification assay as well as quantitative polymerase chain reaction with the assay being negative in all 15 normal skin control samples, giving it a specificity of 100% (95% confidence interval 78.2% to 100.00%) and, a positive predictive value of 100% and negative predictive value of 83.3% (95% confidence interval 72.6%-90.4%) and an accuracy of 91% (95% confidence interval 83.2%-96.1%). Only 23/50 (sensitivity 46%; 95% confidence interval 31.8%-60.7%) slit aspirate samples were positive by microscopic observation [Table 2 and Figure 4b]. Amongst these slit aspirate samples, 34 out of 39 multibacillary samples (sensitivity 87.2%; 95% confidence interval 72.6%-95.7%) and eight out of 11 paucibacillary samples (sensitivity 72.7%; 95% confidence interval 39%-94%) were positive by loop-mediated isothermal amplification assay whereas only 21 multibacillary samples (sensitivity 53.9%; 95% confidence interval 38.6%-68.4%) and only two paucibacillary samples (sensitivity 18.2%; 95% confidence interval 5.1%-47.7%) were found positive by microscopy [Table 2].

Discussion

Early diagnosis of leprosy is critical to control, along with its transmission and prompt treatment, as the nerve damage associated with the disease is permanent and irreversible. At present, diagnosis is based on clinical manifestations followed by the classical confirmatory tests such as histopathology or microscopy; however, these techniques suffer from low sensitivity. The cost-effective isothermal amplification-based techniques have emerged to substitute expensive molecular methods.¹⁸ Amongst these isothermal amplification fulfills all the requisites of the WHO-ASSURED criteria – affordable, sensitive, specific, user-friendly, robust and reliable, equipment-free and deliverable to those in need. Till date, loop-mediated isothermal amplification has been



Figure 4: (a) Visual detection of amplified loop-mediated isothermal amplification products in clinical samples. Tissue samples of the, (b) Visual detection of amplified loop-mediated isothermal amplification products in clinical samples. Slit aspirates samples

Table 2: Sensitivity of loop-mediated isothermal amplification assay in tissue punch samples and slit aspirate samples

Assays	Sensitivity %, 95% CI			
	Multibacillary patients (<i>n</i> =39)	Paucibacillary patients (<i>n</i> =11)		
Tissue punch samples (<i>n</i> =50)				
qPCR	100% (n=39/39), 1-100%	100% (<i>n</i> =11/11), 74-100%		
LAMP	94.9% (<i>n</i> =37/39), 82.7-99.4%	90.9% (<i>n</i> =10/11), 58.7-99.8%		
Histopathology	66.6% (<i>n</i> =26/39), 51-79.4%	45.5% (<i>n</i> =5/11), 21.3-72%		
Slit aspirate samples (<i>n</i> =50)				
qPCR	87.2% (<i>n</i> =34/39), 72.6-95.7%	72.7% (n=8/11), 39-94%		
LAMP	87.2% (<i>n</i> =34/39), 72.6-95.7%	72.7% (n=8/11), 39-94%		
Microscopy	53.9% (<i>n</i> =21/39), 38.6-68.4%	18.2% (n=2/11), 5.1-47.7%		
LAMP: loop-mediated isothermal amplification, CI: confidence interval, qPCR:				

quantitative polymerase chain reaction

successfully applied in the diagnosis of various diseases such as leishmaniasis,²² tuberculosis,²³ malaria,²⁴ dengue,²⁵ chikungunya,²⁶ African trypanosomiasis²⁷ and many more. Herein, we report a highly sensitive loop-mediated isothermal amplification assay specific for leprosy and its application on tissue and slit aspirate samples of leprosy patients. The loopmediated isothermal amplification primers were designed against the *RLEP* gene, a highly conserved repetitive sequence with high copy number (n = 37).¹⁰ The developed assay was highly sensitive showing an analytical sensitivity of 100 fg that is equivalent to approximately 30 *M. leprae* organisms. An earlier attempt on developing leprosy specific loop-mediated isothermal amplification assay showed lower sensitivity (~50 *M. leprae* organism) without demonstrating its utility in clinical samples.²⁸

The loop-mediated isothermal amplification assay for leprosy showed 100% specificity when tested on a range of organisms causing co-endemic diseases such as tuberculosis, leishmaniasis and malaria and other bacterial infections. In the present study, we used closed-tube technology that gave better sensitivity, clarity in the visualization of the results and minimized the risk of cross-contamination associated with conventional loop-mediated isothermal amplification thereby increasing its applicability in the field. The closed-tube loopmediated isothermal amplification assays, either by using waxdye capsules^{29,30} or adding dye in tube cap at the beginning of reaction^{22,31} has been used widely to avoid cross-contamination. Our loop-mediated isothermal amplification assay for leprosy showed an overall sensitivity of 94% using tissue samples which were significantly higher than the routinely used histopathology (62%). The assay also showed a distinct advantage in detecting paucibacillary cases with 90.9% sensitivity in comparison with histopathology (45% sensitivity).

In our study, we attempted loop-mediated isothermal amplification assay using slit aspirates to minimize invasive procedure of sample collection. Skin slit aspirate is far less invasive than tissue biopsy as it does not require anesthesia, suturing, long healing time and leaves no mark after healing. Among different sampling techniques, slit aspirate sampling has the advantages of ease of collection, storage and transportation and most importantly can be effective in case-contact studies, a requisite to control leprosy transmission.³² The loop-mediated isothermal amplification assay was as effective as a quantitative polymerase chain reaction for detection of M. leprae (sensitivity 84%) using slit aspirate sample while being markedly superior to microscopy (sensitivity 46%). In fact, the loop-mediated isothermal amplification assay performed better than the reported polymerase chain reactionbased molecular tests that exhibited a sensitivity of approximately 76%.14,17 Furthermore, 8 out of 11 (72.7%) slit aspirates from paucibacillary patients tested positive for M. leprae as compared to two out of 11 (18.2%) by routine microscopy. Thus, based on the results observed in paucibacillary patients, our loop-mediated isothermal amplification assay can indeed be advantageous for detecting M. leprae in the early stage of the disease which could be crucial in controlling disease transmission.

Limitations

The paucibacillary patients constituted only 22% of the total cases enrolled in the study. Our loop-mediated isothermal amplification assay needs to be extended to a larger number of paucibacillary cases to establish its utility in early detection of the disease.

Conclusion

Loop-mediated isothermal amplification was shown to be a simple and rapid method for the laboratory identification of *M. leprae* in clinical samples. Our loop-mediated isothermal amplification assay has advantages over presently available diagnostics in terms of sensitivity and specificity and can undoubtedly be volunteered as an efficient tool for diagnosis of leprosy as it is simpler, faster, cost-effective and easy to perform with the potential of field applicability.

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Declaration of patient consent

The authors certify that they have obtained all appropriate patient consent.

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Conflicts of interest

There are no conflicts of interest.

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