# Role of molecular approaches to distinguish post kala-azar dermal leishmaniasis from leprosy: A case study

## Sir,

Visceral leishmaniasis, or kala-azar, is caused by the kinetoplastid parasite Leishmania donovani and its dermal sequel, post kala-azar dermal leishmaniasis appears after apparent cure from visceral leishmaniasis. Post kala-azar dermal leishmaniasis (PKDL) in South Asia occurs in approximately 10% of cases with a lag period following visceral leishmaniasis that ranges from 2 to 10 years.<sup>1</sup> PKDL presents with either macular lesions or a combination of papules, nodules and macules, termed polymorphic. In visceral leishmaniasis endemic areas, cases of leprosy or Hansen's disease caused by Mycobacterium leprae can present with similar macular or nodular lesions, along with nerve thickening and hypoesthesia, thus posing a diagnostic dilemma.<sup>2</sup> As per the World Health Organization algorithm for diagnosis of post kala-azar dermal leishmaniasis, the criteria include a history of visceral leishmaniasis,



Figure 1a: A 36-year old male with an isolated macular hypoesthetic patch on lower right back

rK39 immunochromatographic strip test positivity, presence of amastigotes in slit skin smears, direct agglutination test and/or ELISA positivity (https://apps.who.int/iris/bitstream/ handle/10665/78608/9789241505215\_eng.pdf). In view of the clinical dilemma, molecular approaches like quantitative real-time polymerase chain reaction and Sanger sequencing are employed to reach a conclusive diagnosis.

During an active field survey and medical camp conducted in Harirampur block, South Dinajpur district, West Bengal, a 36-year-old male presented with a macular hypoesthetic patch on the lower right side of back. The lesion was present for eight years along with ulnar nerve thickening [Figures 1a and b]. At presentation, the patient had no fever, hepatosplenomegaly or lymphadenopathy, but gave a history of visceral leishmaniasis 32 years ago for which he received sodium antimony gluconate.



Figure 1b: Magnified view of the macular lesion on lower right back

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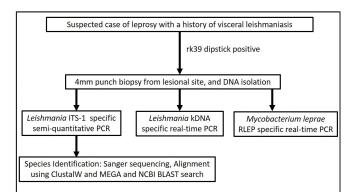
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The rK39 dipstick positive DNA was and was isolated from a 4 mm skin biopsy. Primers for Leishmania specific Internal Transcribed Spacer-1 gene (LITSR 5'-CTGGATCATTTTCCGATG-3'. 5'-TGATACCACTTATCGCACTT-3')3 L5.8S and kinetoplastid minicircle gene (kDNA F: 5'-CTTTTCTggTCCTCCgggTAgg-3'; **k**DNA 5'-CCACCCggCCCTATTTTACACCAA-3') R: along with a 5'FAM tagged probe (5'-6FAM-TTTTCgCAgAACgCCCCTACCCgC-BBQ were used to perform polymerase chain reaction and quantitative realtime polymerase chain reaction, respectively.<sup>4</sup> The Internal Transcribed Spacer-1 polymerase chain reaction 320 base pair (bp) product was visualised by 2% agarose gel electrophoresis, while the cycle threshold for the suspected case was 17.42. The Leishmania donovani WHO reference strain MHOM/IN/80/DD8 was a positive control and was positive by Internal Transcribed Spacer-1 polymerase chain reaction and quantitative polymerase chain reaction, cycle threshold being 20.19. The non-template control was undetermined. For quantification of parasite load, a standard curve was generated by serially diluting DNA isolated from  $1 \times 10^7$  Leishmania donovani parasites and a quantitative polymerase chain reaction was performed.3

A *Mycobacterium leprae*-specific repetitive element (RLEP) based Taqman quantitative polymerase chain reaction was similarly undertaken using *Mycobacterium leprae* specific primers (RLEP-F: 5'-gCAgTATCgTgTTAgTgAA-3'; RLEP-R: 5'-CgCTAgAAggTTgCCgTATg-3') and 5'FAM tagged probe (5'-6FAM-CgCCgACggCCggATCATCgA-BBQ) with a *Mycobacterium leprae* TN strain being the positive control.<sup>5</sup> Cycle threshold value of the positive control was 27.99 and for the case, it was undetermined, confirming the patient was negative for leprosy [Figure 2].

To confirm the Leishmania species, cycle sequencing and capillary electrophoresis was undertaken. Briefly, the chromatogram obtained was converted into a fast alignment file and analysed using Molecular evolutionary genetics analysis. Reference sequences of Internal Transcribed Spacer-1 of different species of Leishmania were retrieved from GenBank (http://www.ncbi.nlm.nih.gov/GenBank/index.html) and aligned with the patient sequence via ClustalW and Molecular evolutionary genetics analysis software version 11.0 (Accession numbers KJ465104.1, KJ465105.1, MN244152.1 for Leishmania donovani, MH347947.1 for Leishmania tropica and KJ002553.1 for Leishmania. major). The patient's sequence aligned with Leishmania donovani sequences. Furthermore, a BLAST search showed a 99.9% match with Leishmania donovani (Accession number KJ465105.1). (https://blast. ncbi.nlm.nih.gov/Blast.cgi?PROGRAM=blastn&PAGE\_ TYPE=BlastSearch&LINK LOC=blasthome, Figure 3). The patient's sequence was submitted to GenBank (Accession number ON817264).



**Figure 2:** A proposed algorithm to distinguish post kala-azar dermal leishmaniasis from leprosy. BLAST, Basic local alignment search tool; ITS-1, Internal transcribed spacer-1; kDNA, Kinetoplast DNA; MEGA, Molecular evolutionary genetics analysis software; NCBI, National Center for Biotechnology Information; PCR, Polymerase chain reaction; RLEP, M. leprae-specific repetitive element

Score 492 bits(266)		Expect 1e-134	Identities 272/275(99%)	Gaps 1/275(0%)	Strand Plus/Plus	
492 00	13(200)	16 154	2/2/2/3(33/0)	1/2/3(070)	Flus/Flus	
Query	9	CTTCGGGGGGGRACCTA	TGTATATATATATGTAGG			8
Sbjct	1	CTTCGGGGAG-ACCTA	TGTATATATATATGTAGG			9
Query	69	TGTMCTCAAAATTTGC	AGTaaaaaaaGGCCGATC	GACGTTATAACGCACCG	CTATACAA 1	28
Sbjct	60	TGTACTCAAAATTTGC	AGTAAAAAAAGGCCGATC	GACGTTATAACGCACCG	CTATACAA 1	19
Query	129	AAGCAAAAATGTCCGT	ТТАТАСААААААТАТАСG	GCGTTTCGGTTTTTGGC	GGGTGGGT 1	88
Sbjct	120	AAGCAAAAATGTCCGT	ТТАТАСААААААТАТАСG	GCGTTTCGGTTTTTGGC	GGGTGGGT 1	79
Query	189	GCGTGTGTGGATAACG	GCTCACATAACGTGTCGC	GATGGATGACTTGGCTT	CTATTTCG 2	48
Sbjct	180	GCGTGTGTGGGATAACG	GCTCACATAACGTGTCGC	GATGGATGACTTGGCTT	CTATTTCG 2	39
Query	249	TTGAAGAACGCAGTAA	AGTGCGATAAGTGGTATC	A 283		
Sbjct	240	TTGAAGAACGCAGTAA	AGTGCGATAAGTGGTATC	A 274		

Figure 3: BLAST alignment of sequence sourced from the patient (Accession number ON817264) and Leishmania donovani sequence having accession number KJ465105.1 (98.9% match). BLAST, Basic local alignment search tool

Post kala-azar dermal leishmaniasis cases form a relatively neglected component of the ongoing visceral leishmaniasis elimination programme, but being potent reservoirs for disease transmission, need urgent attention. The identification of macular post kala-azar dermal leishmaniasis is challenging as their hypopigmented lesions mimic leprosy.<sup>6</sup> As this patient presented with a single hypopigmented hypoesthetic patch and ulnar thickening, strongly suggestive of leprosy along with a prior history of visceral leishmaniasis, quantitative polymerase chain reaction for post kala-azar dermal leishmaniasis and leprosy was necessary. The positivity of rK39, Internal Transcribed Spacer-1 polymerase chain reaction and quantitative polymerase chain reactionR confirmed Leishmania infection. Additionally, as the quantitative polymerase chain reaction of Mycobacterium leprae specific repetitive element was negative, it corroborated this was a case of post kala-azar dermal leishmaniasis, masquerading as leprosy. As it was a single lesion, the causative species could have been Leishmania donovani, Leishmania major or Leishmania tropica. Therefore, it was necessary to define the causative species which was facilitated by Sanger

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sequencing, and confirmed the causative organism was *Leishmania donovani*.<sup>7</sup>

Due to the considerable overlap between macular post kala-azar dermal leishmaniasis and leprosy, and the prolonged period of therapy necessary, multiple molecular approaches were applied for confirming the causative organism. Nerve inflammation and peripheral nerve involvement are symptoms suggestive of leprosy, but neural involvement has been reported in Sudanese post kala-azar dermal leishmaniasis<sup>8</sup> as also in a case from Bihar.<sup>9</sup> Additionally, symptoms of leprosy can also overlap with psoriasis, pityriasis versicolor, granuloma annulare, syphilis, vitiligo and sarcoidosis.<sup>10</sup> Taken together, suspected leprosy or post kala-azar dermal leishmaniasis cases should ideally be tested for other diseases to ensure effective case management.

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#### Ethical approval

The study was approved by the Institutional Ethics Committee.

### Declaration of patient consent

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

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#### Conflict of interest

There are no conflicts of interest.

# Sutopa Roy, Madhurima Roy, Shankha Nath<sup>1</sup>, Surya Jyati Chaudhuri<sup>2</sup>, Manab K Ghosh<sup>3</sup>, Souvik Mukherjee<sup>1</sup>, Mitali Chatterjee

Department of Pharmacology, Institute of Postgraduate Medical Education and Research, Kolkata, <sup>1</sup>Department of Biotechnology, National Institute of Biomedical Genomics, Kalyani, <sup>2</sup>Department of Microbiology, Sarat Chandra Chattopadhyay Government Medical College and Hospital, Uluberia, Howrah, <sup>3</sup>Department of Tropical Medicine, School of Tropical Medicine, Kolkata, West Bengal, India.

> Corresponding author: Prof. Mitali Chatterjee, Department of Pharmacology, Institute of Postgraduate Medical Education and Research, 244 B, AJC Bose Road, Kolkata 700 020, West Bengal, India. ilatimc@gmail.com

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