Evidence for *Mycobacterium leprae* Drug Resistance in a Large Cohort of Leprous Neuropathy Patients from India

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Abstract. Resistance to anti-leprosy drugs is on the rise. Several studies have documented resistance to rifampicin, dapsone, and ofloxacin in patients with leprosy. We looked for point mutations within the folP1, rpoB, and gyrA gene regions of the *Mycobacterium leprae* genome predominantly in the neural form of leprosy. DNA samples from 77 nerve tissue samples were polymerase chain reaction (PCR)-amplified for *M leprae* DNA and sequenced for drug resistance–determining regions of genes rpoB, folP1, and gyrA. The mean age at presentation and onset was 38.2 ± 13.4 (range 14-71) years and 34.9 ± 12.6 years (range 10-63) years, respectively. The majority had borderline tuberculoid leprosy (53 [68.8%]). Mutations associated with resistance were identified in 6/77 (7.8%) specimens. Mutations seen were those associated with resistance to rifampicin, ofloxacin, and dapsone. All the six patients were drug-naive. The clinical and pathological manifestations in this group did not differ from the drug-sensitive group. This study highlights the occurrence of resistance to the standard multidrug therapy and ofloxacin in leprosy. Among the entire cohort, 1/77 (1.3%) showed resistance to rifampicin, 2/77 (2.6%) to dapsone, and 5/77 (6.4%) to ofloxacin. Six new patients showing infection by mutant strains indicated the emergence of primary resistance. Resistance to ofloxacin could be due to frequent use of quinolones for many bacterial infections. The results of the study indicate the need for development of a robust and strict surveillance system for detecting drug resistance in leprosy in India.

INTRODUCTION

Leprosy is one of the leading causes of peripheral neuropathy, particularly in low-resource or developing countries. Leprosy affects the sensory, motor, and autonomic nerve functions, leading to prominent sensory loss and muscle weakness.^{1–3} Despite three decades of effective treatment with multidrug therapy (MDT), leprosy remains a public health problem. Recently, there has been an increase in relapsing cases, failures in MDT treatment, and the emergence of drug-resistant strains of *Mycobacterium leprae*, which could undermine existing leprosy control measures.

Standard treatment for leprosy is an MDT with dapsone, rifampicin, and clofazimine.⁴ Resistance to dapsone and rifampicin is reported from many leprosy-endemic regions of the world.^{5–7} Although prevalence of leprosy is declining it remains a global problem with potential for resurgence in many areas as borders open and worldwide travel is fast.⁸

The emergence of MDR organisms for any infectious disease is a public health concern. Currently, recommended control measures for treating leprosy with MDT are designed to prevent the spread of dapsone-resistant *M. leprae* strains. In the current study, patients with leprous neuropathy were studied for resistance to rifampicin, dapsone, and ofloxacin by PCR sequencing of the drug resistance–determining regions (DRDR) of the genes rpoB, folP1, and gyrA. Clinical, histopathological, and treatment status in the drug-resistant and drug-sensitive groups has been reported.

MATERIALS AND METHODS

Study subjects. This ambispective cohort study was conducted at the National Institute of Mental Health and Neuro Sciences (NIMHANS), Bangalore, located in the southern part of India. It is a national quaternary care center for all neurological disorders. National Institute of Mental Health and Neuro Sciences is not a center identified specifically for leprosy patients, but patients with predominantly pure neuritic form of leprosy or leprous neuropathy with unidentified skin lesions attend the outpatient clinics. In this study, spanning over a period of 12 years (2007 to 2018), a total of 83 nerve samples of leprosy patients were obtained. A final number of 77 samples were selected for reporting. We were able to procure 44 preserved specimens of nerve tissue from histopathologically characterized patients of leprosy, after careful scrutiny of all neuropathology records with diagnosis of leprosy. The remaining 33 patients were prospectively recruited from 2015, and fresh nerve tissue was taken up for PCR study.

Ethics statement. All patients included in the study were adults. As part of the hospital requirement, written informed consent was obtained from all the patients for diagnostic biochemical tests and biopsy procedure. Ethics approval from the "NIMHANS Ethics Committee" (Institutional Review Board) was obtained to collect all clinical, biochemical, and histopathological characteristics from the hospital medical records and process all stored nerve tissue samples of leprosy patients for PCR study and sequencing for DRDR in rpoB, foIP1, gyrA—the genes responsible for rifampicin, dapsone, and ofloxacin drug resistance, respectively.

Detection of mutations by PCR targeting rpoB, folP, and gyrA. All DNA samples isolated from preserved nerve tissue and fresh nerve tissue stored in RNAlater solution were transferred to Stanley Browne Laboratory, The Leprosy Mission Trust India in Delhi. Nerve tissue from paraffin blocks were analyzed with FFPE DNA (M/s. Analytik Jena AG, Jena, Germany) kit method for DNA extraction. *Mycobacterium leprae*–specific DNA was amplified by PCR and sequenced for the aforementioned genes. Of the 83 specimens tested, the *M. leprae* DNA from 77

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samples was PCR-amplified (18/77 were fresh nerve tissue samples preserved in RNAlater solution).

PCR methodology. PCR-based gene amplification was performed using primers according to the guidelines of WHO "Global Surveillance of Drug Resistance in Leprosy 2008" for detection of mutation in rpoB, gyrA, and folP1 genes in M. leprae genome.9 The PCR mixture contained 12.5 µL Hot Start Tag polymerase PCR Master Mix (2 ×) (Qiagen, Hilden, Germany), 1.25 µL forward primer and reverse primer at a final concentration of 0.5 μ M and 5 μ L of template DNA from the processed sample. The final volume of the reaction mix was made up to 25 µL with nuclease-free water. Primer sequences used in this study were as follows: folP primers: folPF CTTGATCCTGACGATGCTGT; folPR CCACCAGACACATCGTTGAC, rpoB primers: rpoBF GTCGA GGCGATCACGCCGCA; rpoBR CGACAATGAACCGATCA GAC, and gyrA primers: gyrAF ATGGTCTCAAACCGGTA CATC; gyrAR TACCCGGCGAACCGAAATTG. The reaction was cycled 40 times at 94°C, 60°C, and 70°C for 1 minute each; preceded by initial denaturation at 95°C for 15 minutes; and ended by final extension at 72°C for 10 minutes. Each reaction setup contained one negative and one positive control. After detection of the PCR product on a 2% agarose gel, amplicons were excised from the gel and later purified

using the Qiagen Gel Extraction Kit. The PCR products were sent for commercial sequencing (at Eurofin Technologies Pvt. Limited, Bangalore, India). Sequence data were analyzed using MEGA version 5.1 (http://megasoftware.net/).

RESULTS

Patients came from different states of India and also Nepal (Figure 1). Based on the clinical and nerve histopathology findings, 83 patients of leprosy with neuropathy were selected for the study. DNA could be isolated from all the 83 nerve tissue specimens. *Mycobacterium leprae*–specific DNA was amplified by PCR from 77 of 83 (93%) specimens, and these were included for the final analysis. The biopsied nerves were superficial radial nerve = 33 (42.9%); dorsal cutaneous branch of ulnar nerve = 18 (23.4%); sural nerve = 17 (22.1%); and superficial peroneal nerve = 9 (11.7%).

The six patients' DNA samples not amplified by PCR were also reported to have leprosy by histopathology (healed leprosy = 2, lepra reaction = 1, and borderline tuberculoid [BT] = 3 [one showed acid fast bacilli]). Among the 77 patients, the clinical diagnoses were leprous neuropathy in 70/77 (91%), mononeuritis multiplex in four (6.3%), vasculitic neuropathy in

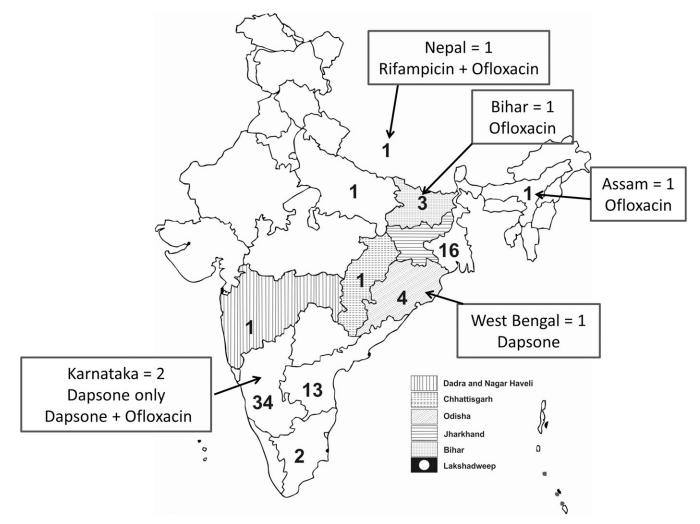


FIGURE 1. Schematic diagram of map of India and Nepal. Embedded numbers represent number of leprosy patients from each state in India and from Nepal. Data in boxes indicate the number of resistant patients and the drug. The shaded regions represent the areas endemic for leprosy.

two (3.2%), and distal symmetric sensory motor neuropathy in one (1.6%). Patients were from different states of the country and neighboring nation of Nepal. Family history of leprosy was reported in 5/77 (6.5%) patients. For the entire group, the mean age at the onset of disease, age at presentation, and duration of illness were 34.9 ± 12.6 (range 10-63) years, 38.2 ± 13.4 (range 14-71) years, and 3.4 years (range 1 month to 55 years), respectively. The M: F ratio was 70:7 (10:1). The patient with 55 years duration is a 71-year-old man with recurrence of neuropathic symptoms. This duration was purely based on the patient's version of his prior chronic neuropathic illness at 16 years of age, when he was prescribed anti-leprosy drugs for 6 months. His past medical records were not available for verification. The majority of patients (43/77, 56.0%) presented between 20 and 40 years of age (60.0%), and 46/77 (60.0%) also had onset between 20 and 40 years. Among the 77 specimens, 72 had a definitive diagnosis by histopathology. One patient demonstrated mild axonopathy in the thickened but unaffected superficial radial nerve. However, fine needle aspiration of the ulnar nerve abscess revealed M. leprae DNA. The six patients with resistance had a mean age at onset of 30.0 \pm 16.8 (range 13–61) years, mean age at presentation of 33.0 \pm 17.0 (range 14–61) years, and mean duration of illness of $3.1 \pm$ 4.4 (range 1–12) years. Five of these six patients had BT and one had borderline lepromatous (BL) leprosy by histopathology (Table 1). Among the 71 remaining patients with no evidence of resistance, the mean age at onset of symptoms was 35.1 ± 13.0 (range 10–63) years, age at presentation was 38.3 ± 13.8 (range 14-71) years, and duration of illness was 4.7 ± 7.4 years (1 month-2 years). The histopathological diagnosis was as follows: BT = 48, tuberculoid (TT) = 8, BL = 7, mid borderline (BB) = 3, healed/fibrosed = 4, and axonopathy = 1. The place of origin and salient features of the six patients with resistance are summarized in Table 1. All except one patient from Bihar hailed from nonendemic regions for leprosy in India. The demographics and clinical and histopathological features of the group with resistance did not differ from those of the remaining larger drugsensitive group. No unusual features were encountered in the drug-resistant group.

Resistance mutations were identified in 6/77 (7.8%) specimens: rifampicin (Leu436GIn) combined with ofloxacin (Ala91Pro) in one, dapsone (Pro55Leu) combined with ofloxacin (Ala91Val) in one, only dapsone (Pro55Ser) in one, and only ofloxacin in three patients (Gly89Ala, Ala91Pro, and Gly89Cys) (Table 2).

The clinically affected nerve was biopsied in 75 (97.4%) patients. The nerve selection, histopathological findings in the biopsied nerve, and its classification are represented in Table 1. AFB was demonstrated in 11 (14.3%) specimens. Skin lesions were reported in 27 patients (35.0%), but examination revealed lesions in 42 patients (54.5%). In addition to nerve biopsy, skin biopsy from the affected area was performed in 15 patients, and this showed the following: BT = 4, BB = 1, TT = 3, nonspecific inflammation = 5, and no changes = 2. Nerve conduction studies carried out in the 77 patients showed sensory motor demyelinating polyradiculoneuropathy in 76 (98.7%) patients and motor sensory axonopathy in 1 (1.3%) patient.

DISCUSSION

This is a 12-year ambispective cross-sectional observational study conducted at a quaternary center for neurological disorders. The retrospective group had 44 patients, from 2007 to 2015, and the prospective group had 33 patients, from 2016 to 2018. The main aim was to look for mutations in the *folP1*, *rpoB*, and *gyrA* genes in *M. leprae* DNA isolated from 77 biopsied peripheral nerve tissue characterized to have leprosy by histopathology. Both drug-naive and MDT-treated patients were studied, and findings with regard to the resistance pattern for *M. leprae* along with the demographics and histopathological features have been presented. There were six patients identified to have resistance. Importantly, all patients were drug-naive, indicating primary infection with resistant strains. Among these, one patient came from Bihar which is reported to be an endemic region for leprosy.

Although the prevalence of leprosy in India is less than 1/10,000 since 2005,¹⁰ it still accounts for 62% of the total new patients reported worldwide. Emergence of multidrug-resistant *M. leprae* strains is reported in several studies from Southeast Asia and other parts of the developing world. This indicates a serious threat to effective control of leprosy, particularly in South Asia. It is proposed that development of drug-resistant strains of *M. leprae* is particularly due to premature discontinuation of treatment and monotherapy.^{11,12} In India, dapsone monotherapy has been discontinued since 1985. *Mycobacterium leprae* resistance to dapsone was first reported in 1964,¹³ for rifampicin in 1976,¹² and for ofloxacin in 1996 in nude mice with established *M. leprae* infection,¹⁴ and in the year 2000 in a Japanese patient for dapsone, rifampicin, and sparfloxacin.¹⁵

The program for Global Sentinel Surveillance for drug resistance in leprosy was established by the WHO in 2009 to monitor drug resistance among patients of relapse.⁹ Nine cases of dapsone and one case of rifampicin resistance were detected among 72 leprosy relapse patients.¹⁶ In contrast, in our cohort, six (7.8%) new drug-naive leprous neuropathy patients demonstrated resistance to either one or more anti-leprosy drugs. Five of the six patients had BT and one had BL leprosy. Two of our BT patients had resistance to dapsone: one to dapsone alone and the other to dapsone and ofloxacin. The other three BT patients had resistance to rifampicin and ofloxacin and one each to ofloxacin only, and the BL patient to ofloxacin. Thus, the findings suggest that the infection was caused by drugresistant M. leprae, an indication of emergence of primary resistance. The clinical features, histopathological characteristics, type of leprosy, and treatment regime did not differ between the drug-resistant and drug-sensitive groups.

Point mutations in *M. leprae* genome responsible for dapsone and rifampicin resistance have been identified in several studies, and this has facilitated the development of new tools for monitoring drug resistance.^{17,18} In a large study from Tamil Nadu, India, 214 clinically suspected patients of relapse/drug resistance were investigated for MDT susceptibility by mouse foot pad inoculation. Authors identified 15 (16%) secondary dapsone-resistant strains.¹⁹ In comparison, two patients with primary resistance to dapsone were identified.

Mycobacterial resistance to rifampicin correlates with changes in the structure of the subunit of the DNA-dependent RNA polymerase, primarily because of missense mutations of the rpoB gene.²⁰ Our single patient with resistance to rifampicin also had a missense mutation but with dual resistance to ofloxacin also.

In a study from Kolkata, India, skin biopsies from 50 patients with a relapse of leprosy were studied for rifampicin resistance.

Variables	Case 1	Case 2	Case 3	Case 4	Case 5	Case 6
Place of origin	Darjeeling, West Bengal	Nepal	Kadapa, Andhra Pradesh	Dharbhanga, Bihar	Dhubri, Assam	Chikkaballapur, Karnataka
Year of evaluation	2008	2013	2014	2015	2017	2018
Endemic region	No	No	No	Yes	No	No
Follow-up in	Lost	Lost	Deficits stable. MDT	Lost	Deficits stable.	Deficits stable.
December 2018			for 12 months 5		On MDT for 12	Stopped MDT
			years ago. No		months	after the 12-month
			active disease			course
Gender	М	М	M	F	М	М
Age at presentation (years)	45	22	61	14	32	24
Age at onset (years)	33	20	60	13	31	22
Duration of illness (months)	144	24	18	12	24	48
Affected limb(s)	Lowerlimbs	Lower limbs	Upper limbs	Right upper limb	Lower limbs	Right upper limb
Histopathology diagnosis	BT	BT	BL	BT	BT	BT
Biopsied nerve	DCUN	Superficial	DCUN	DCUN	SRN	SRN
·		, peroneal				
		nerve				
Histopathology findings						
Plasma cells	-	_	Yes	-	-	-
Histiocytes	Yes	Yes	Yes	-	-	-
Lymphocytes	Yes	Yes	Yes	Yes	Yes	Yes
Perineurial thickening	Yes	Yes	Yes	Yes	Yes	Yes
Endoneurial thickening	Yes	Yes	Yes	Yes	Yes	Yes
Perivascular inflammation	Yes	Yes	Yes	Yes	Yes	Yes
Giant cells	-	-	-	-	Yes	-
Foam cells	-	Yes	-	-	-	-
Caseous necrosis	-	Yes	-	-	-	-
AFB	No	No	Yes	No	No	No
Skin lesions	Yes	Yes	No	No	Yes	Yes
Hypopigmented patches	Yes	-	-	-	Yes	Yes
Nonhealing ulcers	Yes	Yes	-	-	-	-
Mutations in drug resistance-	-determining re	egion				
Dapsone	Pro55Ser	-	-	-	-	Pro55Lue
Rifampicin	-	Lue436Gln	-	-	-	-
Ofloxacin	-	Ala91Pro	Ala91Pro	Gly89Ala	Gly89Cys	Ala91Val
Prior drug therapy	None	None	None	None	None	None

TABLE 1 Salient demographics, and clinical and histopathological characteristics in the six leprosy patients with resistant mutations

BT = borderline tuberculoid; DCUN = dorsal cutaneous branch of ulnar nerve; SRN = superficial radial nerve.

Two patients who had completed 2 years of MB-MDT showed a missense mutation in the CAC codon.²¹ Our case with resistance to rifampicin also had the same mutation. In a report from Brazil, *M. leprae* DNA from nerve tissue was tested for mutations in rpoB, folP1, gyrA, and gyrB genes. Mutations in the rpoB and folP1 genes were detected, suggesting multidrug resistance. Seventy percent of the *M. leprae* mutations were isolated from relapsed patients. No resistance to ofloxacin was detected.²² One of the mutations (Pro55Leu) for dapsone resistance was the same as seen in this study. Thus, the mutations identified for dapsone resistance seem to be common.

In a study carried out in Vietnam, skin smears/skin biopsy samples from 423 leprosy patients showed 19 mutations in 187 folP1 samples, but none related to drug resistance in 163 rpoB and 147 gyrA gene samples was noted. Only two of the 19 were new cases. In comparison, all the six (7.8%) patients with resistance were new cases in the present study group.

Patients with relapse and those who had received dapsone monotherapy in the past have shown high mutation rates (78%), compared with patients with relapse who had previously received MDT (33%).¹¹ In contrast to that report, the findings in this study indicate that resistant strains are causing infection in our group. The first case in the present study had undergone nerve biopsy in 2007, and it means that resistant strains have caused leprosy several years ago. No follow-up information on this patient is available.

In a study from Louisiana, among 39 patients, one case each with dapsone alone and dapsone- and rifampin-resistant *M. leprae* was detected,^{23,24} and both were drug naive. Dapsone-resistant patients in this study also had distinct mutations with primary resistance.

In the 1980s, the WHO recommended dapsone, rifampicin, and clofazimine as the primary agents for MDT.⁴ In this series, none of the patients exhibiting drug resistance had received any anti-leprosy drug(s). There were 28 (36.4%) patients who had received MDT in the past for varying periods before nerve biopsy, and 49 (64.6%) were drug-naive at the time of biopsy.

In a previous study carried out on slit skin smears of 250 samples, from India, MDR *M. leprae* was detected from relapsed leprosy patients. Seven had showed mutations for rifampicin and dapsone, seven for dapsone and ofloxacin, and one for rifampicin and ofloxacin resistance.²⁵ By contrast, the present cohort was studied for *M. leprae* DNA in nerve tissue and showed dual resistance in two patients similar to the aforementioned report and single drug resistance in the remaining four. However, in our comparatively smaller sample size, a case with dual rifampicin and ofloxacin resistance was identified, which otherwise is a rare occurrence.

In a worldwide study conducted in 19 countries, including India, 1932 patients were studied for mutations in rpoB, folP1, and gyrA genes. Dual resistance to rifampicin and dapsone and to dapsone and ofloxacin was identified, but none to a

TABLE 2 Mutation pattern of drug resistance for multidrug therapy

SI.No	Drugs	Leprosy type	Mutation	Position	Change in nucleotide
1	Rifampicin and ofloxacin	BT	Leu436GIn and Ala91Pro	279 bp (2275228–2275506) 2275524 , 225 bp (7499–7682)	$CTG \rightarrow CAG$ (T is replaced by A) GCA \rightarrow CCA (G is replaced by C)
2	Dapsone and ofloxacin	BT	Pro55Leu and Ala91Val	254 bp (296800–297025) 296859 , 225 bp (7499–7682)	$CCC \rightarrow CTC$ (C is replaced by T) GCA \rightarrow GTA (C is replaced by T)
3	Dapsone	BT	Pro55Ser	254 bp (296800-297025) 296859	$\mathbf{C}CC \rightarrow \mathbf{T}CC$ (C is replaced by T)
4	Ofloxacin	BT	Gly89Ala	225 bp (7499–7682)	$GGC \rightarrow GCC$ (G is replaced by C)
5	Ofloxacin	BT	Gly89Cys	225 bp (7499–7682)	G GC \rightarrow T GC (G is replaced by T)
6	Ofloxacin	BL	Ala91Pro	225 bp (7499–7682)	$GCA \rightarrow GTA$ (C is replaced by T)

BT = borderline tuberculoid. Bold indicates position of mutation and change in nucleotide.

combination of rifampicin and ofloxacin. In our cohort, a single case of rifampicin resistance (1/77, 1.3%), who also had resistance to ofloxacin, was identified. A similar case was reported in only one earlier study.²⁵

In the current cohort, the overall resistance rate among 77 patients was 7.8%, similar to the report in the large multinational study. However, individually, it was less for rifampicin (1/77, 1.3%) and dapsone (2/77, 2.6%), except for ofloxacin (4/77, 5.2%) which was similar. The factor(s) responsible for primary resistance despite MDT practice is of immense concern and, thus, an intense surveillance mechanism should be established under the leprosy control program. More studies on Indian leprosy patients should be carried out to identify the burden of resistance. There are a limited number of studies assessing the importance of identification of M. leprae DNA predominantly in the neural form and pure neural leprosy (PNL).²⁶⁻³⁰ This form of leprosy presents as peripheral neuropathy without skin lesions, and in India and Nepal, it accounts for 3.9-18.0% of all diagnosed patients.³¹⁻³⁵ In this study, there were 40 (52.0%) patients qualified for PNL, which is much higher when compared with previous reports, and this could be due to a referral bias. PCR identified M. leprae DNA in all these patients.

With the findings of resistance in the cohort, it may be valuable to subject tissue samples for PCR testing from all suspected leprosy patients and also to look for the presence of drug resistance. In patients with TT leprosy, histopathological confirmation is not certain as AFB is uncommonly seen and nonspecific inflammation is more frequent. In such situations, subjecting the nerve tissue for PCR for *Mycobacterium lepra* identification may be considered. Furthermore, in all pure neuritic leprosy patients identification of *M. lepra* DNA by PCR will confirm the diagnosis as mislabeling leprosy as chronic inflammatory demyelinating neuropathy, neurosarcoidosis, and vasculitis could unnecessarily lead to long-term incorrect therapy and, thus, adverse effects of immunomodulation.

In conclusion, the current series presents the pattern of drug resistance identified in a large cohort of leprosy patients with primarily neural form of leprosy. For the first time, sequencing for the DRDR for MDT drugs was performed on DNA isolated from nerve tissue in India. In this study, 6/77 patients with drug resistance and two patients with dual resistance were identified. PCR testing for identification of *M. leprae* DNA is a simple, fast, and affordable method of confirming leprosy and should be taken up based on the type of leprosy. Furthermore, the isolated *M. leprae* DNA should be subjected to determination of drug resistance so that treatment strategies can be modified accordingly and also the spread of resistant strains can be curtailed.

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