

DETECTION OF *Mycobacterium leprae* DNA FOR 36kDa PROTEIN IN URINE FROM LEPROSY PATIENTS: A PRELIMINARY REPORT

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SUMMARY

We have searched for *Mycobacterium leprae* DNA for 36kDa protein in urine using a *M. leprae* specific PCR technique. A limited number of 16 patients (of which 11 belonged to lepromatous leprosy and five to tuberculoid leprosy) and eight healthy individuals were included for the present study. The number of urine samples positive by PCR were 36.4% (4/11) in lepromatous patients and 40% (2/5) in tuberculoid patients. None of the samples from healthy individuals was positive. To our knowledge, the results indicate, for the first time, the presence of *M. leprae* DNA in urine from leprosy patients. Another important finding obtained out of the study is that amongst treated patients 66.6% (4/6) were positive whereas amongst untreated only 20% (2/10) were positive. From the present indicative data it appears that treatment improves the PCR results with urine as a sample. Thus, the approach could prove to be useful for monitoring the treatment response of individual patients and needs to be further evaluated with a large number of patients.

KEYWORDS: Urine; Leprosy; *Mycobacterium leprae*; DNA; PCR; Diagnosis

Leprosy is a chronic infectious disease caused by *Mycobacterium leprae*. The disease manifests in various forms resulting in a spectrum⁸. At one pole of the spectrum lies tuberculoid leprosy (TT) while at the other one lies lepromatous leprosy (LL). Between these two polar types various borderline forms (i.e. BT-borderline tuberculoid, BB-mid-borderline, BL-borderline lepromatous leprosy) have been described. In tuberculoid leprosy there is a localized skin and/or nerve lesion(s), with a low bacterial load and strong cell mediated immunity. In contrast, lepromatous leprosy has generalized lesions with a high bacterial load and strong humoral immunity. Immunological examination of urine from leprosy patients has shown that samples from a proportion of leprosy patients are positive for antigens and antibodies in the urine^{4,5,6}. Though, there are reports documenting mycobacterial DNA in urine from patients suffering from infectious diseases^{7,10}, to our knowledge, there is hardly any information, demonstrating *M. leprae* DNA in urine from leprosy patients. With skin biopsies from leprosy patients a specific PCR for detection of DNA for 36kDa antigen has already been published^{1,2} in the literature. In this pilot study, we have made an attempt to detect *M. leprae* DNA for 36kDa protein in urine from leprosy patients.

Sixteen active leprosy patients (who attended the out patient department at Central JALMA Institute for leprosy, Agra) and eight healthy individuals were included for the study. Patients were diagnosed using clinical criteria and classified according to RIDLEY & JOPLING⁸ scale. None of the patients had any symptom indicating the defective

renal function. Of the 16 patients, six were on anti leprosy treatment. Various relevant clinical features have been depicted in Table 1.

After obtaining the informed consent from all the individuals, they were requested to produce urine. Ten ml of the urine sample was centrifuged at 12,074 g for 20 min at 4 °C. The supernatant was discarded and the pellet was suspended in 500 µl of 10 mM Tris-HCl supplemented with 1 mM ethylene diamine tetra-acetic acid (TE, pH 8.0). If needed, samples were stored at -20 °C till they were used.

The suspension of urine sediment (430 µl) in TE was boiled for 10 minutes. It was then followed by sudden cooling and treatment with 50 µl lysozyme (20 mg/ml) for two hrs at 37 °C. Next, 5 µl of proteinase-K (10 mg/ml) and 56 µl of 10% sodium dodecyl sulphate (SDS) were mixed and incubated for one hour at 65 °C. Then, 80 µl of 5 M sodium chloride (NaCl) and 64 µl cetyl tri-methylammonium bromide (CTAB; 20mg/ml) were added to the mixture and incubated for 30 min at 65 °C. The suspension thus obtained was treated with equal volume of chloroform-iso-alcohol (1:24) solution, vortexed and centrifuged at 12,074 g for five minutes. Supernatant was mixed with iso-propanol (0.6 times by volume) in a separate tube. The contents of the tube were kept at -20 °C overnight. Later on, the tube was centrifuged at 12,074 g for 15 min, sediment in the tube was mixed after discarding the supernatant. Seventy percent ethanol (150 µl) was then added to the mixture and the tube was centrifuged for 7 minutes at 12,074 g. Finally, after drying the ethanol

Table 1
Details of patients analyzed by PCR

Patient	Disease duration (Months)	Treatment duration (Months)	Bacterial Index (BI)	Type of disease	PCR result
1	6	Untreated	0	BT	+
2	12	Untreated	0	BT	-
3	1.5	Untreated	3	LL	-
4	36	Untreated	5	LL	-
5	24	5	4	LL	+
6	24	Untreated	3	BL	-
7	72	1.5	5	LL	-
8	3	3	4	LL	+
9	18	4	3	BL	+
10	12	2	3	LL	-
11	36	Untreated	3.5	LL	-
12	18	Untreated	4	LL	+
13	6	6	0	BT	+
14	36	Untreated	0	BL	-
15	12	Untreated	0	BT	-
16	3	Untreated	0	BT	-

from the sample, 20 µl of TE buffer was added to the contents in the tube and preserved at -20 °C till use. While performing this procedure all precautions were taken to prevent cross contamination.

DNA extracted from samples were subjected to PCR following the methods, with slight modification, described previously by HARTSKEERL *et al.*² and DE WIT *et al.*¹ where the target for amplification was 530 bp sequence on the gene encoding the 36kDa antigen. The method included both, positive and negative controls. The denaturation, annealing and extension steps were carried out at 94 °C for two minutes, at 60 °C for two minutes and at 72 °C for three minutes respectively. The amplicon obtained out of this procedure was analyzed by agarose gel electrophoresis incorporated with 0.05 µl/ml of ethidium bromide (0.5 mg/ml). On completion of the run the separated PCR products were transferred to positively charged nylon membrane. The transferred products were then probed with a digoxigenin labeled 530 bp fragment specific for DNA encoding for 36kDa antigen. All steps including labeling of the probe, hybridization and detection employing anti-digoxigenin antibody conjugated with alkaline phosphatase using the substrate 5-bromo-4-chloro-3-indolyl phosphate and nitroblue tetrazolium salt were performed as described by the manufacturer (Roche Diagnostics, Mannheim, Germany). A specimen was considered positive when a 530 bp band was revealed by both agarose gel electrophoresis and subsequent hybridization.

The overall positivity rate (Table 2) with samples from leprosy patients was 37.5% (6/16). None of the healthy controls was detected to be positive. The group-wise positivity was 36.4% (4/11) with samples from lepromatous patients and 40 percent (2/5) with samples from borderline tuberculoid patients. In BT group of patients the cell mediated immune response is known to be comparatively better than LL/BL type of patients⁸. Probably, there may be continuous damage (at slow rate) of *M. leprae* in BT patients resulting in more DNA release in the blood and thereby more excretion in the urine leading to better positivity, though

Table 2
Performance of PCR for detection of *M. leprae* DNA in urine

Leprosy status	Positive	Negative	Positivity (%)
Lepromatous	4/11	7/11	36.4
Tuberculoid	2/5	3/5	40
All	6/16	10/16	37.5
Healthy	0/8	8/8	0
Untreated	2/10	8/10	20
Treated	4/6	2/6	66.7

insignificant, when compared to BL/LL patients. On the other hand, with LL/BL patients the situation is reverse. Further, no association between PCR status and clinical features including bacterial indices were found. To conclude, detecting urinary DNA for 36kDa protein does not appear to be a helpful tool for diagnosis of *M. leprae* infection.

In leprosy, efforts have been made in the past to demonstrate *M. leprae* DNA in a variety of clinical samples such as skin, lymph, blood, nasal secretions, hair bulbs and fresh biopsy⁹. To our knowledge, present is the first report regarding demonstration of *M. leprae* DNA in urine from leprosy patients. The appearance of *M. leprae* DNA in urine from leprosy patients does not seem to be surprising as prevalence of antigens and antibodies in urine from leprosy patients is already known^{4,5,6}. However, mechanism of getting DNA and even antigen and antibody is not known as yet.

One interesting observation obtained out of the present study is that amongst treated patients 66.7% (4/6) were positive whereas only 20% (2/10) samples were positive from untreated group. Surprisingly, all the patients having taken three or more than three months treatment were positive by PCR. This indicative data points out that probably treatment

affects the PCR results by enhancing the release of DNA in the urine. It is known that due to treatment *M. leprae* is damaged and then eliminated from the host tissues³. During this process, various bacterial components including bacterial DNA might be released in the host tissues and DNA thus released might get way in the urine. Hence, detecting DNA in urine appears to be worthwhile to explore further on large number of samples for establishing its utility in monitoring the response to chemotherapy in leprosy patients.

RESUMO

Deteção do DNA do *Mycobacterium leprae* para proteína 36 kDa na urina de pacientes com hanseníase: relato preliminar

Pesquisamos o DNA do *Mycobacterium leprae* para proteína 36 kDa na urina usando a técnica do PCR específica para *M. leprae*. Um número limitado de 16 pacientes (dos quais 11 tinham hanseníase multibacilar e cinco hanseníase paucibacilar) e oito indivíduos saudáveis foram incluídos neste estudo. O número de amostras de urina positivas pelo PCR foi de 36,4% (4/11) em pacientes com hanseníase multibacilar e 40% (2/5) em pacientes com hanseníase paucibacilar. Nenhuma das amostras de indivíduos saudáveis foi positiva. Até onde chega o nosso conhecimento, os resultados indicam, pela primeira vez, a presença de DNA do *M. leprae* na urina de pacientes com hanseníase. Outro fato importante obtido através do exame é que entre os pacientes tratados 66.6% (4/6) eram positivos enquanto entre os não tratados somente 20% (2/10) foram positivos. Pelos presentes dados indicativos parece que o tratamento melhora os resultados do PCR em amostra de urina. Assim, o acesso a estes dados prova ser útil no monitoramento da resposta ao tratamento de pacientes individuais e precisa ser melhor avaliado com um grande número de pacientes.

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