

Value of Polymerase Chain Reaction (PCR)
In Diagnosis and Follow-up of Paucibacillary Leprosy
Patients

Thesis

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List of Abbreviations

AFB	: Acid Fast Bacilli
APC	: Antigen Presenting Cells
BB	: Borderline Leprosy
BI	: Bacterial Index
BL	: Borderline Lepromatous Leprosy
BT	: Borderline Tuberculoid Leprosy
CMI	: Cell mediated Immunity
DC	: Dendritic Cells
DNA	: Deoxyribo Nucleic Acid
DTH	: Delayed Type Hypersensitivity
ENL	: Erythema Nodosum Leprosum
H&E	: Haematoxyline – Eiosine
HD	: Hansen's Disease
IFN-γ	: Interferon Gamma
IL – γ	: Interleukin-γ
IL – γ	: Interleukin – γ
LL	: Lepromatous Leprosy
LC-PCR	: Light Cycler Polymerase Chain Reaction
LIB	: Logarithmic Index of Biopsies
M. leprae	: Mycobacterium Leprae
MB	: Multibacillary
MDT	: Multi Drug Therapy

MI	: Morphological Index
NAPCS	: New – Antigen Presenting Cells
PB	: Paucibacillary
PCR	: Polymerase Chain Reaction
PGL-I	: Phenolic Glycolipid-one
QPCR	: Quantitative Polymerase Chain Reaction
RT. PCR	: Real- Time Polymerase Chain Reaction
SGPSI	: Stocking- Glove Pattern Sensory Impairment
TLRs	: Toll Like Receptors
Th – 1	: T .lymphocytes Helper Cells
TNF- &	: Tumour Necrosis Factor Alpha
TT	: Tuberculoid Leprosy
VC	: Virchow Cells
WHO	: World Health Organization

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Introduction

Leprosy, a chronic granulomatous disease which primarily affects the peripheral nervous system and skin, has a worldwide distribution; the total number of cases is estimated to be around ๐๓๔,๐๐๐. India accounts for ๖๔% of leprosy prevalence and ๙๖% of the new cases detected worldwide (*Dayal et al, ๑๙๙๗*).

Leprosy is characterized by two polar forms, a paucibacillary tuberculoid and a multibacillary lepromatous (LL) form. Between these, three intermediate forms are described: the borderline tuberculoid (BT), borderline borderline (BB), and borderline lepromatous (BL) forms. Another rare form is the pure neural leprosy (PNL), which is characterized by the uniqueness of nerve lesions and is difficult to diagnose because skin lesions and acid-fast bacilli (AFB) in slit smears are absent (*Riedly and Jopling 19๙๗*).

The diagnosis of paucibacillary leprosy is primarily clinical. Anesthetic skin lesions with or without thickened peripheral nerves are virtually pathognomonic of paucibacillary leprosy, a full thickness skin biopsy from an anesthetic lesion showing granuloma and lymphocytic infiltration of nerves essentially confirms the diagnosis. Acid fast bacilli (AFBs) are rarely found in patients with indeterminate or tuberculoid disease. In the established form of the disease, the diagnosis is relatively simple. However, the established

forms are decreasing in number due to widespread use of multi drug therapy (MDT) and early disease is being reported more frequently (*Buhrer-Sekula et al, 1997*).

In early suspected cases, routine histopathological examination by Hematoxylin and Eosin and Fite-Faraco staining of sections can confirm the diagnosis in only about 30% of such early cases, because of paucity of acid fast bacilli and absence of infiltration inside the dermal nerves . Instead, they show non-specific histopathology in the form of chronic inflammatory cell infiltrate at various locations, which is not specific for leprosy alone (*Ramu et al, 1997*).

The diagnosis of these early cases requires additional methods as demonstrating nucleic acid sequences specific to the pathogen by e.g. using in-situ hybridization and amplification by in situ polymerase chain reaction (*Jardim et al, 1997*). This technique allows cellular localization of low-level of nucleic acid within the tissue and cells and can aid in a significant way to augment the sensitivity of histopathological diagnosis, and provide a solid base for follow up of these patients (*Komminoth et al, 1997*).

Aim of the work

The aim of this work is to evaluate the technique of Real-time polymerase chain reaction (RT-PCR) in the diagnosis of paucibacillary leprosy i.e tuberculoid leprosy, borderline tuberculoid and the pure neural leprosy. Another aim is to assess the value of (PCR) technique in the follow up of the patients for the efficacy of the multidrug therapy (MDT).

Introduction

Leprosy is a chronic granulomatous disease principally affecting the skin and peripheral nerves caused by the infection with the obligate intracellular *Mycobacterium leprae*. (*scollard et al., 2007*)

Leprosy is best understood as two conjoined diseases. The first is a chronic mycobacterial infection that elicits an extraordinary range of cellular immune response in humans. The second is a peripheral neuropathy that is initiated by the infection and the accompanying immunological events. (*Sunaina et al., 2014*)

Epidemiology

The prevalence of leprosy fell by 90 percent in the last two decades because patients completing a course of multiple-drug therapy(MDT) have been considered to be cured, but the incidence of the disease, which varies in direct proportion to case-finding efforts, has yet to be reduced convincingly. (*WHO, 2010* .)

The highest new case detection rates are in India, Brazil, Democratic Republic of Congo, Tanzania, Nepal, Mozambique, Madagascar, Angola and the Central African Republic. The disease burden in India alone represents 64% of all new cases worldwide. (*WHO, 2010* .)

At the beginning of 2010, the registered prevalence of leprosy globally was 211,903 cases and the number of new cases detected during 2009 was 244,796 cases. (WHO, 2010) (Table 1)

Table (1): Prevalence of leprosy and number of new cases detected, by WHO region, beginning of 2010. (WHO, 2010)

WHO region	No of cases registered (prevalence rate*) Beginning of 2010	No of new cases detected (case-detection rate**) 2009
African	30,647 (0,40)	28,930 (3,70)
Americas	43,370 (0,49)	40,474 (4,08)
South-East Asia	120,406 (0,68)	166,110 (9,39)
Eastern Mediterranean	8,490 (0,10)	40,29 (0,70)
Western Pacific	8,630 (0,00)	0,243 (0,29)
Total	211,903	244,796

* Prevalence rate = number of cases / population

** Case-detection rate = number of cases / population

In all populations studied, lepromatous disease is more common in men than in women by a 2:1 ratio. If leprosy is common, the tuberculoid forms dominate; if leprosy is uncommon, the lepromatous forms predominate. Leprosy is said to be a disease of rural incidence, but of urban prevalence. The median age of onset is less in tuberculoid than in lepromatous patients, but in both groups,

the median age of onset being less than 30 years old. However, old age does not protect. (WHO, 1994)

In Egypt, the highest new case detection rates are present in the Upper Egypt Governorates (Qena, Sohag, Aswan and Assiut, respectively). (WHO, 1994)

Table (4): Leprosy situation in Egypt, beginning of 1990 (WHO, 1994).

Registered prevalence	No. of new cases detected 1994	No. of new cases of multibacillary leprosy	No. of new female cases	No. of new cases among children	No. of new cases with grade 2 disabilities	No. of relapses 1994
912	700	616	260	48	42	0

Mode of transmission:

The route of infection of leprosy is unproven, but current evidence favors respiratory transmission: evidence for congenital and percutaneous transmission has been presented, but examples are rare. (Truman, 1994)

The incubation period for tuberculoid leprosy is up to 10 years and for lepromatous disease may be 20 years or longer. (Irgens, 1994) Untreated multibacillary (MB) patients are probably the most important source of Mycobacterium leprae transmission. It is estimated that household contacts of MB patients have a relative

risk of developing leprosy that is 5- to 10-fold greater than that of the general population. However, in many areas, the number of MB patients is very small, and they may not represent the most important source of infection. **(Cole et al., 1997)** There is increasing evidence that subclinical transmission may occur, since even in countries where leprosy is highly endemic, for many patients, no history of close contact with a leprosy patient can be established. **(International Leprosy Association Technical Forum, 1998)**

As *Mycobacterium leprae* can persist and possibly proliferate in the environment in association with certain plants and animals, it is conceivable that infection may result through prolonged or repeated exposure to an environmental source containing viable bacilli. This is difficult to investigate experimentally because *M. leprae* cannot be cultivated in vitro and evidence can only be obtained indirectly through epidemiological studies. **(Desikan and Sreevatsa, 1998)**

Genetics of susceptibility:

A twin study has provided compelling evidence that both genetic and environmental factors are important in determining disease susceptibility and disease expression. The study demonstrated higher concordance rates for leprosy among monozygotic compared with dizygotic twins. **(Chakravarti and**

Vogel, 1993) Various genes and regions in the human genome have been linked to or associated with susceptibility to leprosy per se or with a particular type of leprosy. These are not all reproducible in different populations, which may be unsurprising (*Ciechanover, 1997*).

Mira et al. (1998) have identified a certain locus within the gene PARK2\ PACRG which is located on chromosome 2 and was identified to be associated with overall susceptibility of human populations to *M. leprae* (*Mira et al., 1998*). This is the first example of the use of positional cloning to identify a human gene associated with susceptibility to an infectious disease (*Skamene, 1998*). In their initial association scan of a linkage peak identified through a genome scan of a Vietnamese patient population, the investigators identified a locus within this gene that was highly associated with leprosy (leprosy per se), regardless of the subtype of this disease. These results were confirmed by a second analysis of Brazilian families with one or more persons affected by leprosy. The specific locus is a promoter region of PARK2 and a coregulated gene, PACRG, located on chromosome 2q35-q37. PARKIN was named for its association with an early-onset form of Parkinson's disease, and the finding that a locus within this gene is also associated with susceptibility to leprosy is very unexpected (*Liu et al., 1997*).