

Original Research Article

Identification of Immunotopes against *Mycobacterium leprae* as Immune Targets Using PhDTm- 12mer Phage Display Peptide Library

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Abstract

Purpose: To determine the surface epitopes of *Mycobacterium leprae* (*M. leprae*) and evaluate their efficacy in the production of anti-*M. leprae* antibodies in an animal model.

Methods: Blood samples were obtained from 34 patients suffering from lepromatous leprosy. Antibodies were obtained from the samples, semi-purified and used to coat the wells of ELISA microplate, and M13 random-peptides library was added to the wells. After four rounds of panning, three clones were isolated and their peptide mimotopes were sequenced. Western blot was used to evaluate the interaction of the isolated mimotopes.

Results: Three selective clones were tested by direct enzyme-linked immunosorbent assay (ELISA) and western blot. anti-*leprae* antibodies in various dilutions and were found to be serological active. Sequencing of the isolated peptides showed identities between the two clones that were able to successfully induce anti-*Leprae* humoral response in mice.

Conclusion: The findings indicate that the isolated peptides can potentially be used for early diagnosis. However, further research is required to improve their potency as new vaccines against leprosy.

Keywords: Bacteriophage, Vaccine, Leprosy, *Mycobacterium leprae*, Random peptide phage display library

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INTRODUCTION

Leprosy is a chronic granulomatous bacterial infection with *M. leprae* that principally affects the skin and the peripheral nervous system [1-2]. In endemic areas, the incidence of leprosy is estimated to be about 4 to 6 cases per thousand. The lipid-rich cell wall of pathogenic bacteria are likely responsible for their resistance and enable

bacteria to escape from host defense mechanism.

Leprosy treatment is difficult due to slow growth rate of the bacteria and long incubation period (average of 3-10 years). Additionally, multiple relapse and the emergence of drug-resistant strains have made the treatment more challenging. Such clinical prognosis correlates with the level of cell-mediated immunity (CMI)

against *M. leprae*. Antibody responses are great in lepromatous patients, indicating that humoral immunity does not contribute to the host defense [6].

On the other hand, full assessment of *M. leprae* antigenic structure is not possible due to lack of bacterial growth in culture media that has major contribution in the absence of suitable vaccine against leprosy [4]. The World Health Organization's effort to eliminate leprosy by 2000 was based on important advances achieved in anti-mycobacterial therapy in the 1980s. Although over the last decade, the dramatic reduction in prevalence of leprosy has been reported in most of the endemic countries, but its incidence rate is also high [3,5]. In this context, the IDEAL consortium Initiative for Diagnostic and Epidemiological Assays for Leprosy was established in 2004 and represents an international task-force that brings together researchers from endemic and non-endemic countries and aiming at the concerted development of diagnostic tests for early diagnosis of leprosy [5]. Identification of molecular markers improves epidemiological knowledge and transmission of leprosy that leads to intervention designing for prevention of the disease [3,7].

Display systems based on bacteriophages are widely used to assess different proteins and antibodies developed for identification of pathogenic bacteria [8-10]. A phage library that expresses random short peptide sequences on the surface of filamentous phage represents one of the biotechnological tools that could be helpful in amino acid sequence identification of various antigenic epitopes [11-13]. Panning the peptide phage library against a specific antibody results in isolation and identification of desired epitopes and their amino acid sequence and also determining the nucleotide sequence of genes expressing peptides on the phage surface [11,13,14].

In this research, the M13 random peptide phage library was used to detect the surface antigen epitopes of leprosy bacterium [15]. In addition to high efficiency, phage display systems are easy to work with and require no advanced techniques, expensive equipment or materials [16]. Isolated phage-peptides can be used for induction of immune response without use of adjuvants [17,18]. M13 phage is a filamentous bacteriophage and is one of the most used display systems [16]. One of the important and unique features of the vectors derived from the M13

phage is their potential for expressing short random peptide sequences on one of their coat proteins [16].

In this study, specific anti-*M. leprae* antibodies isolated from patients were used in panning against peptide phage library. This method can help to identify the functional epitope sequences and may contribute to development of a diagnostic assay or vaccines against leprosy [15,19].

EXPERIMENTAL

Library preparation

The Ph.D.TM-12 Phage Display Peptide Library is based on a combinative library of random peptides fused on a minor coat protein (pIII) of M13 phage [20,21]. The peptide is linked to the pIII coat protein by a spacer (Gly-Gly-Gly-Ser). The library consists of approximately 2.8×10^9 unique members, amplified once to yield approximately 70 copies of each sequence in 10 μ l of the supplied phage.

Preparation of leprosy antiserum

Antiserum was prepared from blood samples of 34 leprosy patients in Lepromatous stage that were received from Center for Research and Training in Skin Disease and Leprosy. Plasma was separated from the blood clot after centrifugation at 1000 g at 4 °C. Serum samples of all 34 patients were pooled together and ammonium sulfate precipitation (final concentration 45 %) was used to increase the concentration of antibodies. After addition of sodium azide, the precipitated serum was distributed in 500 μ L aliquots and kept at -20 °C. Protein concentration of samples were determined by the Bradford method [22].

Phage titration

The phage library was titrated before panning. Serial dilutions of phage were prepared in LB broth (1 % Yeast extract, 1 % Tryptone and 1 % NaCl). Dilutions were added to *E. coli* ER2738 and incubated at room temperature for 5 minutes. The bacteria were mixed with Top agar (1 % Yeast extract, 1 % Tryptone, 1 % NaCl and 0.75 % Agar) and plated on LB agar containing X-Gal (20 μ g/mL) and Isopropyl thiogalactose (IPTG) (1 mM). The plates were incubated at 37 °C for 16 h and numbers of colonies were counted to obtain phage pfu/mL.

Screening

The ELISA microplate wells were coated with 15 µg of leprosy antiserum in 150 µL volume of 100 mM sodium bicarbonate buffer (pH 8) and incubated overnight at 4 °C. The wells were washed using TBS (50 mM Tris-Cl, 150 mM NaCl, pH 7.5) and blocked using 200 µL of 100 mM sodium bicarbonate buffer containing 3 % BSA and incubated at 37 °C for 1 h. Wells were then washed six times using TBS containing 0.05 % Tween 20 (TBST). A total of 4×10^{10} Phages were diluted in 1 ml of TBST buffer and 100 µL was added to each well. The microplate was incubated at room temperature for 1 h. The supernatant was discarded and the wells were washed 10 times with Tris-Buffered Saline and Tween 20 (TBST) buffer. For eluting the bound phages, 100 µL of glycine and BSA buffer (100 mM HCl-Glycine, pH 2.2) was added to the wells and the plate was incubated at room temperature for 15 min. Finally, 15 µL of 1 M Tris buffer (pH 9.1) was added to each well for neutralization of glycine buffer.

Phage multiplication

One microliter of isolated phages was added to 200 mL of fresh bacterial suspensions at mid log phase and incubated for 4.5 h at 37 °C. It was centrifuged at 10000 g for 10 min at 4 °C. The supernatant was saved and mixed with 20 % (v/v) PEG/NaCl solution (16 % Polyethylene glycol in 2.5 M NaCl) and incubated on ice for 16 h. The Phage/PEG/NaCl solution was centrifuged at 15000 g for 20 min. The phage sediment was resuspended in 1 mL TBS solution and stored at 4 °C for further application. To obtain the highest affinity binders, panning procedure ("Screening" and "Phage multiplication" steps) was repeated three or more times. Eluted phages from previous round were used for each panning round.

Clone selection

The phages after the fourth panning round were used for selecting specific clones with high affinity obtained leprosy antiserum.

For selection of specific clone, 100 µg of patient anti-serum, normal anti-serum and BSA (5 mg/ml) blocking buffer was separately added to the wells in duplicate and the plate was incubated 16 h at 4 °C.

Serial dilution was prepared from phages purified from all selected clones. For each clone, 1×10^{12} phage were diluted in a volume of 200 ml and

then diluted 12 times by a factor less than 4, until number of phages reached 2×10^5 pfu/mL.

After washing and blocking the wells, 100 µl of the diluted phages were added to each well and the plate was incubated at 37 °C for 1 h. Wells were washed and 200 µl of the anti-M13 antibody (New England Biolabs, UK) diluted in the blocking buffer (1:5000) was added to each well. After 1 h incubation at 37 °C, the wells were washed 6 times with TBST buffer. Finally, 15 min after addition of the ABTS substrate (2, 2'-Azinobis [3-ethylbenzothiazoline-6-sulfonic acid]-diammonium salt) (Sigma, Germany) dissolved in sodium citrate buffer the resulting color was read at 405-415 nm by ELISA reader. The clones with high absorbance compared with the normal serum and BSA were considered as positive clones.

Western blot analysis

To confirm phage expression of the peptides, 10 % SDS-PAGE was used and the proteins transferred onto nitrocellulose membrane using a semi-dry transfer apparatus (ATTO Co. Japan), and thereafter incubated by polyclonal Anti-*M. leprae* MLMA-LAM (Rabbit antiserum, BEI Resources). The secondary antibody (HRP-conjugated rabbit anti-IgG antibody, Abcam), diluted in the blocking buffer, was added. After washing the membrane with TBST for 15 min, detection was performed using DAB (diaminobenzidine) as a substrate [13,20].

Animal tests

Six weeks old Balb/c mice (15-18 g) were used to evaluate the immunogenicity and specificity of the peptide in induction of anti-*M. leprae* antibodies.

The experiment included 6 groups each of 5 mice. The mice in groups 1, 2 and 3 were injected with 200 µl of 1×10^{13} cfu /mL of clone 6, 9 and 10, respectively. Groups 4 and 5 were considered as positive controls and were respectively injected with 100 µg of Lepromin (ML) and Mycobacterium extract (M) in 200 µL PBS. Group 6 was the negative control and each mouse in the group was injected with the wild M13 bacteriophage in the same amount as groups 1 to 3. All injections were carried out intraperitoneally. First booster was injected after 3 weeks and the rest of injections were done with 2 week intervals. The immunization process was assayed by indirect ELISA. *M. leprae* extract was used to coat the ELSA microplate wells. Sera obtained from the mice were used as primary antibody and anti-mouse IgG antibody

conjugated with HRP (AbCam, UK) was used as a secondary antibody.

Sequence determination

To determine the genetic sequence of the peptides, 100 μ L of positive sample mixed in TE buffer (1×10^{13} cfu/mL) was sequenced by primers of 12-mer Peptides of Phage Display Library kit (Biolab, England).

Statistical analysis

One-way analysis of variance (ANOVA) was used to assess differences in the binding of phage-peptide to the test serum, control serum and BSA. The data are presented as mean \pm standard deviation and compared using Duncan's multiple range tests ($p < 0.01$).

RESULTS

Clone selection

The clones were selected from the 4th round. Study of optical absorptions showed that only 3

clones namely; 1, 6, and 9, had notable binding difference between the test serum, control serum and BSA, and showed the highest affinity toward leprosy antiserum. Results show that most clones do not show significant binding differences between the control groups, the test serum and BSA. However, significant difference was observed in comparison of the control groups and test serum in three clones 1, 6 and 9 at 1×10^{12} , 2.2×10^8 and 2×10^5 dilutions.

Positive clones

The size of pIII coat protein was about 45 kDa and can be seen in 45 to 65 kDa range. There were several specific bands within that range that were not observed in the positive and negative controls. These bands are not observed in clone 6. Indeed, clone 6 could not elicit specific anti-*M. leprae* antibody response in the animal tests. The lighter bands could be as a result of protein degradation during the extraction process or cross-reaction. Finally, clones 1 and 9 were accepted as positive (Figure 1).

Table 1: Analysis of ELISA results performed on test and control sera, and BSA at a dilution of 2×10^5 cfu/mL

Subject	Source of variant	Mean square	Sum of squares	df	F _c
Patient Serum	Between Groups (Treatment)	1.279	12.793	10	171**
	Within Groups(Error)	0.007	0.164	22	
Normal Serum	Between Groups (Treatment)	0.004	0.041	10	1.641
	Within Groups(Error)	0.003	0.055	22	
BSA	Between Groups (Treatment)	0.002	0.020	10	1.902
	Within Groups(Error)	0.001	0.023	22	

*Significant at 1% probability

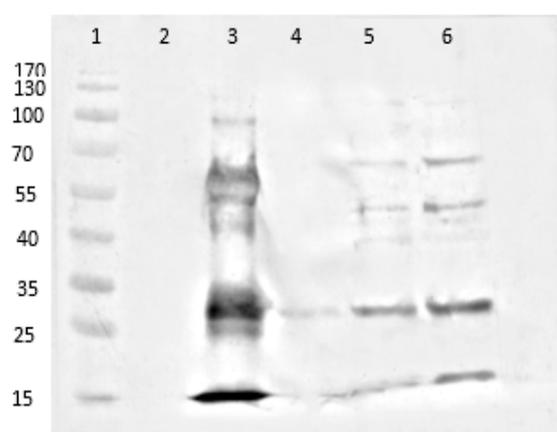


Fig 1: Western blot analysis on selected clones. Lane 1: Standard protein Marker. Lane 2: Protein extracted from wild type M13 bacteriophage. Lane 3: cell extracts of *Mycobacterium Leprae*. Lanes 4, 5 and 6: Protein extracts from phage clones 6, 9 and 1, respectively

Animals

Clone 6 showed immunogenicity only in one round; however, clones 1 and 9 produced specific anti-*M. leprae* immunity in all repeats Table 2. Groups 5 and 6 which were immunized with M and ML form respectively, were used as positive controls and were able to induce high immunity. Group 6 is the negative control group injected with wild type M13 showed no specific immune response against *M. leprae* antigens. Following the results of immunogenicity, the clones positive were sequenced. Nucleic acid sequences of clones 1 and 9 are shown in Table 3.

Phage-peptides (1, 6 and 9), Wild type M13 (10), *M. leprae* extract (M) and Lepromine (ML) were used to evaluate of specific immune response induction in mice. Three repeats were carried out. Clones which were able to induce anti-*M.*

leprae antibodies were sequenced As shown the peptide sequence of both clones are same.

DISCUSSION

In the current study we achieved high affinity peptides (epitopes) that expressed on the surface of phage and are able to bind anti-*M. leprae* specific antibodies. Finally, two clones showed potential immunogenicity in mice. Lack of immunogenicity in the clone 6 may be because of phage mutation or immune system default of mice models that could not able to recognize this peptide as an antigen. However, clones 1 and 9 were able to induce specific antibodies which are able to react with *M. leprae* antigens, thus these mimotope peptides have potential applications as a vaccine against the disease or may be applied as a diagnostic tool for leprosy cases. Despite that the immunogenicity mechanism is unknown, the antibody titer was considerably high. Given the amount of lepromin used in this study, the immune response of the peptides was equal to those of the positive control groups. The obtained immunotopes would thus be suitable for the preparation of sub-unit vaccine candidates against leprosy, or can be used in a more complex form to include cellular immunity response. Furthermore, the monoclonal antibodies achieved against these peptides can be used for development of accurate diagnostic tests.

In this study, 11 plaques were randomly selected and analyzed for expression of foreign peptides, positive single clones sent for sequencing. Results of sequence identification in clones 1 and 9 showed that the both clones have same sequence that could explain the western blotting binding pattern. The specificity of obtained sequence was confirmed by NCBI-Blast software [24,25].

Leprosy diagnosis is mainly based on symptoms and clinical studies and there is no laboratory test to diagnose or predict the disease progression in individuals at risk [26]. There are two forms of disease caused by *M. leprae*. The specific antibody titer is low in tuberculoid paucibacillary leprosy (PB) but strong Th1/IFN- γ response that the role of cellular immunity (CMI) is significant. In the lepromatous multibacillary patients (MB) the antibody titer is high and the cell-mediated immunity (CMI) is low. Thus the (PB) and (MB) laboratory tests for the leprosy diagnosis are based on measuring the cell-mediated immunity and antibodies. However, the laboratory tests should detect (PB) and (MB) forms. These require proteins with specific properties. These proteins can be obtained by recombinant or synthetic techniques [3,7]. Production of Lepromin requires bacterial culture and animal immunization, in our study we were able to produce peptides that can induce the same immune response as Lepromin,

Table 2: Immune response against mimotope peptides in animal models

Experiment repeat ↓	Immunized with →	1	6	9	10	M	ML
1		+	-	+	-	+	+
2		+	+	+	-	+	+
3		+	-	+	-	+	+

Table 3: DNA sequences of selected recombinant phages

Positive clone	Nucleotide acid sequence of epitope	Amino acid sequence of epitope
Clone-1	5'- CTC GAG CAA TGC CAA GAA TCG -3'	LEQCQES
Clone-9	5'- CTC GAG CAA TGC CAA GAA TCG -3'	LEQCQES

thus the purified peptide has the potential to replace lepromin in research and therapy [15,26].

The methods based on IFN- γ and T-cell are not able to detect sub-clinical disease, and patients at the early stages of treatments. Moreover (PB) and (MB) forms cannot be distinguished [3,7]. It is ideal to use the peptides specific to *M. leprae*, recombinant or synthesized proteins in IFN- γ and T-cell methods. It is important to differentiate between the two (PB) and (MB) phases. These require access to the antigens of the bacteria which is complicated with respect to the

difficulties in culturing and antigen isolation from *M. leprae*.

In this study, the phage clones were selected from the phage library panned against antibodies isolated from lepromatous multibacillary (MB) stages. This stage has been selected due to high levels of antibodies against the disease. Also, by accessing specific peptides, patient identification by early diagnosis of infection would be possible by evaluation of specific cellular immunity (CMI) against leprosy [3].

However, given the genetic diversity of people and variety in HLA classes, feasibility of the methods for producing peptides that could result in production of high affinity antibodies that leads to use of diverse and rich pool of antibodies isolated from various people to include genetic diversity [25]. Accordingly, the antibodies used for panning in this research were extracted from a total of 34 patients and pooled to include most HLA varieties.

M13 random phage peptide library was used against antibodies from leprosy patients to isolate the peptides mimicking the immunogenic epitopes of *M. leprae* antigens.

CONCLUSION

This work identifies peptides which bind with high affinity to antibodies against *M. leprae*. Following injection of peptides to the mice model, the monoclonal antibodies produced can facilitate the development of serological diagnostic kits for detection of this bacterium and disease. Furthermore, the peptide may be used for the development of a new generation of vaccines that induce both cellular and humoral anti-*M. leprae* responses.

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