Comparative studies on molecular techniques for detecting *Rickettsia tsutsugamushi* in *Leptotrombidium deliense* and mice

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Abstract

In order to establish the most sensitive and specific method in the detection of *Rickettsia tsutsugamushi* in *Leptotrombidium deliense*, mice and patients, a comparative study on DNA extraction, PCR, hybridization and PCR-hybridization was conducted. The results showed that the sensitivity of PCR-hybridization was 10 times higher than that of PCR and 1000 times higher than that of hybridization. Quick DNA extraction method could take the place of phenol-chloroform method to prepare DNA template when detecting the rickettsiae in mice and patients' blood by PCR. De-inhibitor quick DNA extraction method could improve the detection rate of rickettsiae from chigger mites.

Key words: *Leptotrombidium deliense*, *Rickettsia tsutsugamushi*, PCR, hybridization, PCR-hybridization

Introduction

Scrub typhus is a natural epidemic disease spread by the larval chiggers of *Leptotrombidium* (*Leptotrombidium*) *deliense* (Walch), *L.(L.) arenicola* (Traub) and *L.(L.) fletcheri* Vercammen-Grandjean (Daryl *et al.* 1994). In nature, both mice and chiggers can be the reservoir hosts of *R. tsutsugamushi*. Our focus is on the cycling mechanism of *Rickettsia tsutsugamushi* (Ogata) between chigger mites and mice, and its spread potential. Traditional methods (including animal infection isolating method, immune method and electronmicroscopy in detection of the rickettsiae) have their own disadvantages (Hong *et al.* 1991). Thus molecular techniques for detecting *R. tsutsugamushi* have been developed (Ni *et al.* 1995; Li *et al.* 1994). In order to establish a most sensitive and specific method for the detection of *R. tsutsugamushi* in chiggers and mice, a comparative study on DNA extraction, PCR, hybridization and PCR-hybridization was carried out.

Materials and methods

Mites

A pure line of *L. deliense* routinely cultured in our laboratory was confirmed free of the rickettsial infection by purification tests with different mite stages including larvae, nymphs, preadults and adults.

Chiggers infected with *R. tsutsugamushi* were obtained either by inoculating *R. tsutsugamushi* into the abdomen of normal adults or by allowing mite larvae to bite infected mice, and then culture these mites. Different mite stages were accumulated for use in experiments.

Mice blood and spleen

Pure line BALB/C mice, which was supplied by the Animal Centre of Sun Yat-sen University of Medical Sciences, was infected with *R. tsutsugamushi* by abdominal inoculation, or by bite of rickettsiae-infected larvae, and then cultured for three weeks. When the mice became ill, blood and spleen were collected for experiments.

Blood and spleen were also accumulated from pure line BALB/C mice.

Rickettsial strains

Karp standard strains of *R. tsutsugamushi* were preserved at -70°C in our laboratory since 1991. Karp DNA was extracted by phenol-chloroform method; serial dilutions of 1:10, 1:10², 1:10³, 1:10⁴, 1:10⁵, 1:10⁶ and 1:10⁷ were prepared with TE buffer (10 mM Tris·Cl, pH 8.0; 1 mM EDTA, pH 8.0).

Rickettsia prowazekii (da Rocha-Lima), *R. mooseri* (Monterio), *R. sibericus* (Zdrodovskii) and Gilliam strain of *R. tsutsugamushi* were provided by the Guangdong Sanitation and Anti-epidemic Station. Kato strain of *R. tsutsugamushi* was also preserved by our laboratory. By phenol-chloroform method, DNA of all these samples were prepared.

Patient blood pool

Rickettsiae-infected blood samples from patients were provided by the Department of Paediatrics and the Department of Infectious Diseases, Fushan First Hospital of Guangdong Province. All patients had typical symptoms and signs of scrub typhus characterized by fever and escharosis.

Primers

Four primers were designed from Sta56 gene of *R. tsutsugamushi* and synthesized by Biomedical Inc.(Alambra, USA)

1. 5'-TTGCTAGTGCAATGTCTGC-3';
2. 5'-CCCTGCTGCTGTGCTTGCTGCCG-3';
3. 5'-TCCTCAGCCTACTATAATGCC-3';
4. 5'-CGACAGATGCAACTATTAGGC-3'.

Amplification fragment was 1 kb when PCR was performed with primers 1 and 2. Amplification fragment was 500 bp when PCR was performed with primers 3 and 4.

Molecular techniques

DNA extraction: Three DNA extraction methods were adopted: classic phenol-chloroform method, quick DNA extraction method and chiggers de-inhibition quick DNA extraction method. Quick DNA extraction liquid was prepared by the senior author. Chiggers de-inhibition quick DNA extraction method was as follows: chiggers were triturated in quick DNA extraction liquid and then boiled for 10 minutes. Samples were then put on hybridization membrane. After being dried, the membrane was washed with 1 mol/L NaCl and TE (pH 8.0) to remove inhibitor factors. DNA fixed membrane was employed for PCR.

PCR Assay: PCR was performed by a programmable thermal controller (model PTC-150, MJ Inc.) and the DNA Amplification Reagent Kit (Biomedical Inc.) with Taq polymerase, 1 U per reaction in a total 50 µl reaction volume. The PCR mixture was denatured at 94°C for 30 seconds, annealed at 57°C for 1 minute, and then the chain was extended at 70°C for 2 minutes. This cycle was repeated 35 times and the sample was then held at 4°C overnight. In the case of DNA fixed membrane, it was soaked in PCR reaction mixture for 10 cycles. After that, the membrane was removed and 1 U Taq polymerase was added again, and then PCR was allowed to continue. 2 µl of the first

PCR product (primer 1-2) diluted to 1:10 in TE was used in nest-PCR (primer 3-4). 5 μ l of the product of nest-PCR was electrophoresed on 1% agarose gel with ethidium bromide as the indicator and pBR322/Hinf I as DNA markers. After EB staining, the bands were observed under ultraviolet detector (Furuya 1993).

Hybridization Assay: The probe from Sta56 was marked with digoxigenin by PCR. The temperature for hybridization was 68°C. The hybridization method used follows that in the manual of DIG DNA Labelling and Detection Kit (Boehringer Mannheim Co.).

Detection of PCR products with blotting method (PCR-hybridization): PCR products were put in a water bath at 100°C for 5 minutes. They were then transferred to an ice bath. 2 μ l of the product were dotted and detected.

Results and discussion

Comparison of three DNA extraction methods for detection of R. tsutsugamushi

PCR results of the classic phenol-chloroform method and quick extraction method: DNA from 20 samples of *R. tsutsugamushi* positive mice blood, 15 samples of positive mice spleen and 15 samples of positive patient blood were extracted with the classic phenol-chloroform method and quick extraction method respectively. The positive rate with quick extraction method was slightly lower than with classic method in mice spleen, but there was no significant difference ($P>0.05$). Detection rates with the two methods were the same in patient blood and mice blood, which suggested that quick extraction method could be used to prepare the sample (Table 1). Because the duration of the experiment was shortened, it was possible that the detection of *R. tsutsugamushi* with PCR could be applied in clinical diagnosis.

TABLE 1. The results of PCR after *Rickettsia tsutsugamushi* DNA extraction with phenol-chloroform method and quick extraction method.

Samples	No. of samples	Positive results (%)	
		Phenol-chloroform method	Quick extraction method
Mice blood	20	100.00	100.00
Mice spleen	15	100.00	93.33
Patient blood	15	100.00	100.00

The results of PCR before and after removing inhibitor factors: 30 samples of *R. tsutsugamushi* infected adults, engorged larvae and nymphs of *L. deliense* were selected from laboratory cultures. DNA from each group was extracted with quick extraction method and de-inhibition quick extraction method respectively. It was showed that there was significant difference between the results obtained before and after removal of inhibitory factors (Table 2). The positive rate dramatically increased after removal of inhibitory factors ($P<0.05$), suggesting that the de-inhibition quick extraction method could be used to extract DNA from *R. tsutsugamushi* in chiggers.

TABLE 2. The results of PCR before and after removal of inhibitory factors in *Leptotrombidium deliense*.

Samples	No. of samples	Positive results (%)	
		Before removal	After removal
Mice blood	30	40.00	100.00
Mice spleen	30	26.67	96.67
Patient blood	30	33.33	93.33

DNA extraction from R. tsutsugamushi in chiggers: Classical DNA extraction methods were deemed unsuitable because of great DNA lost and only trace of *R. tsutsugamushi* in chiggers. On the other hand, the purity of DNA for PCR is not very important. Therefore, the quick extraction method has been recommended by some researchers (Daryl *et al.* 1994). Because of the presence of inhibitors of Taq polymerase in chiggers after engorgement, it was necessary to remove them when employed for PCR. In this study we performed de-inhibition quick extraction method with success.

Specificity of PCR, hybridization and PCR-hybridization for detecting R. tsutsugamushi

The detection of three international standard strains of *R. tsutsugamushi* (Karp, Kato and Gilliam), *R. prowazekii*, *R. mooseri*, *R. sibericus* and normal mice blood showed that the three standard strains (Karp, Kato and Gilliam) could be detected by the above three methods, while the other three species of *Rickettsiae* could not. It was suggested that the three methods could be used to detect *R. tsutsugamushi* in samples (data not shown).

FIGURE 1. Sensitivity of PCR for detecting *Rickettsia tsutsugamushi*. Serial dilutes of Karp DNA: A. 1:10, B. 1:10², C. 1:10³, D. 1:10⁴, E. 1:10⁵, F. 1:10⁶, M. pBR322/Hinfl.

Sensitivity of PCR, hybridization and PCR-hybridization to detect R. tsutsugamushi

Seven serial dilutes of Karp strain DNA were detected by the above three methods. It was demonstrated that the highest diluted concentration which could be detected by PCR was $1:10^5$ (Fig. 1), by hybridization was $1:10^3$ (Fig. 2), by PCR-hybridization was $1:10^6$ (Fig. 3). PCR-hybridization was 10 times as sensitive as PCR alone, and 1000 times more than hybridization. Theoretically, the method of DNA agarose electrophoresis and EB staining is not very sensitive. For a fragment of 100 bp, visible bands appear on the gel only after 1010 amplification. The detection limit is 10 ng DNA by agarose electrophoresis. However, the detection limit is 10 pg when hybridization was used in place of agarose electrophoresis. Moreover, some accidental factors may cause nonspecific amplification in PCR, and the size of the amplified bands is similar to that of target bands. As a consequence, false positive result may occur by electrophoresis, but this can be avoided by detection via probes. Therefore, PCR-hybridization can also improve the specificity for detection of the rickettsiae.

FIGURE 2. Sensitivity of hybridization for detecting *Rickettsia tsutsugamushi*. Serial dilutes of Karp DNA: A. $1:10$, B. $1:10^2$, C. $1:10^3$, D. $1:10^4$, E. $1:10^5$, F. $1:10^6$, G. $1:10^7$.

FIGURE 3. Sensitivity of PCR-hybridization to detect *Rickettsia tsutsugamushi*. Serial dilutes of Karp DNA: A. $1:10$, B. $1:10^2$, C. $1:10^3$, D. $1:10^4$, E. $1:10^5$, F. $1:10^6$, G. $1:10^7$.

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