

Original Paper

Isolation of pathogenic leptospire from blood samples of patients by PCR- RFLP method

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Abstract

Background and Objective: Leptospirosis is a worldwide zoonosis that is more common in tropical and semitropical regions and in flat area of Guilan province, northern Iran, caused by pathogenic leptospire. Serotyping by MAT is complicated, expensive and time consuming. Molecular typing can be useful alternative. This study was performed to identify the typing of endemic pathogenic leptospire, isolated in Guilan province, by PCR-RFLP.

Materials and Methods: In this descriptive study, specimens were taken from patients in Razi hospital, Rasht, Iran, from April to August 2008. DNA of all positive cultures were extracted by Phenol-Chloroform method. PCR was performed by using two primer sets: B64-I, B64-II for Kirschner serovars whose PCR product were digested by HinfI for RFLP, and G1, G2 for all other species whose PCR products were digested by DdeI. Band profiles of digested PCR products were compared to band profiles of standard servers to determine species and subspecies.

Results: 65 of totally 107 blood cultures were positive. 56 Samples were amplified by G1, G2 including Interrogans and Borgpeterseni and 9 samples were amplified by B64-I, B64-II including Kirschneri species.

Conclusion: This study showed that the majority of leptospire species are Interrogans and Borgpeterseni. Regarding to several existing problems in Serotyping of leptospire by MAT, PCR-RFLP can be useful for identifying isolating and studying the serovars in different certain species. It seems that PCR-RFLP can be performable for clinical samples for both early diagnosis and characterization.

Keywords: Leptospirosis, Molecular diagnosis, Molecular typing

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