

عنوان مقاله:

Molecular typing of Iranian strains of *Brucella abortus* and *Brucella melitensis* using RAPD-PCR

محل انتشار:

بیستمین کنگره بین المللی میکروب شناسی ایران (سال: 1398)

تعداد صفحات اصل مقاله: 1

نویسندگان:

Ali Mohammad Behroozikhah - *Razi Vaccine and Serum Research Institute. Agricultural, Research Education and Extension Organization (AREEO), Karaj, Iran*

Mohammad Mehdi Feizabadi - *Department of Microbiology, Faculty of Medicine, Tehran University of Medical Science, Tehran, Iran*

Esmaeil Asli - *Razi Vaccine and Serum Research Institute. Agricultural, Research Education and Extension Organization (AREEO), Karaj, Iran*

خلاصه مقاله:

Introduction and Objectives: Molecular Typing of pathogenic bacteria are introduced to identify local and regional strains, tracing the source of infection, probable changes in native strains, and even an evolutionary study of phylogenetic trees. The objective of this study was to identify DNA polymorphisms and genomic fingerprinting of Iranian *Brucella* strains using RAPD-PCR technique. **Materials and Methods:** 101 strains of *Brucella*, including seven strains of reference (S.99, 16M., RB51, S.19 544, Rev.1, H.38) and 94 Iranian field strains from different geographical regions during 1340 to 1382 were investigated. *Brucella* strains were cultured on *Brucella* broth, *Brucella* agar according to recommended methods. And the purity of strains cultivated with standard methods was assured. Then, all species were determined by phage-typing. Total DNA was extracted from the strains using three methods, including a phenol, Chlorophorm and Iso amilalchol. Optimization of RAPD-PCR conditions with nine primers of 13 single individual primers and 13 pair individuals primers with two different strains of *abortus* and *melitensis* were performed separately. The best result was obtained with the AP4 peptide primer with a sequence of 3' -TCA CGC TGC A-A and the primer pair REP- AP4 with 15-mer sequence, 5'-CGC TTA TCG GCC TAC-3' and AP4 for *Brucella* species for the first time. **Results:** 101 strains of *Brucella* strain with AP4 primer were evaluated with optimum RAPD-PCR conditions. The final PCR product in 5.1% electrophoresis gel with ethidium bromide staining revealed 72 different patterns of DNA from 101 strains of *Brucella*. This method, even among strains within a biotype of the phenomenon of polymorphism, was shown to be repeatable. Within the five genotypes, two to four different biotypes were also included. **Conclusion:** RAPD can be useful method to distinguish related bacterial species, Despite the presence of high polymorphism in DNA. in this study, there was a presence of infection among animals of different *Brucella* biotypes, *Brucella abortus*, Biotype III, and *Brucella melitensis* Biotype 1, which are still dominant and indigenous Iranian strains in the past four decades. This technique appears to be a simple, quick and sensitive for the epidemiological investigation of brucellosis.

کلمات کلیدی:

.RAPD-PCR, *Brucella*, Molecular typing, *Brucella* strains

