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Determination of HCV Genotypes in Iranian Isolates by PCR-RFLP

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Abstract:

Hepatitis C is a major cause of liver related morbidity and mortality worldwide and represents a major public health problem. Depending on genomic organization, the virus is divided into six genotypes and a number of subtypes. Different genotypes are seen in different parts of the world. Genotype one is difficult to treat, while genotypes 2 and 3 are easy to treat. Therefore, identification of HCV genotype in patients is necessary to begin and follow up the treatment. In this study, viral genomic materials of 214 patients' sera were detected by nested-RT PCR. Based on genomic differences among different genotypes, the PCR products were digested with proper enzymes and studied by RFLP. Except for one, sequencing of 14 samples, representative of all genotypes, confirmed the results of PCR-RFLP. The results of PCR-RFLP were as follows: 1a (52.88%), 1b (14.01%), 3a (27.57%), 2a (2.1%), 4 (3.44%). This indicates that a high percentage of HCV infected patients in Iran are infected with 1a or 3a genotypes. These findings reveal that the pattern of HCV genotypes in Iran differs from those of other middle-eastern countries.

Keywords:

Genotyping , RFLP

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