

幽门螺杆菌临床分离株的REP-PCR分析

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摘要 应用REP-PCR对来自20例单纯性胃炎和20例消化性溃疡病人的幽门螺杆菌菌株进行基因分型, 并运用SAS软件进行聚类分析。结果显示这些菌株按照基因型被分为两大类, 但两种来源的菌株在两大类中的比例无明显差异($P>0.1$), 提示幽门螺杆菌临床分离株REP-PCR基因型与致病性之间不存在明显相关性。

REP-PCR Analysis of Helicobacter pylori Clinical Strains

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Abstract: Helicobacter pylori strains isolated from 20 gastritis patients and 20 peptic ulcer patients was genotyped by REP-PCR and was clustered with SAS software. These strains are divided into two categories according to their genotype. But the rate of the two sources of strain in the two categories shows no apparent difference ($P>0.1$), indicating that there is no significant close relationship between the genotype and the pathogenicity.

Key words: Helicobacter pylori; REP-PCR; genotype

关键词 [幽门螺杆菌](#) [REP-PCR](#) [基因分型](#)

分类号

扩展功能

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