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从腾冲热海免培养获得的5个16 S rDNA克隆分析

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Analytical study on five 16 S rDNA clones that come from hot spring in Tengchong with a culture-independent approach

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全文: PDF (509 KB) HTML (KB) 输出: BibTeX | EndNote (RIS) 背景资料

摘要 用免培养法(culture-independent approach)从腾冲大滚锅高温热泉中获得的16SrDNA,经克隆筛选,测定了其中5个克隆的16SrDNA插入片段的近全序列.构建的系统发育树分析结果表明,5个克隆中有2个是*Bacillus*属,1个是*Hydrogenobacter*属,1个是*Pseudomonas*属,还有1个在*Thermodesulfobacteriaceae*科,介于*Geothermobacterium*属和*Thermodesulfobacteria*属之间.

关键词: 热泉 免培养法 16SrDNA 克隆

Abstract: Five almost completely 16 S rDNA sequences were obtained from Dagunguo of Tengchong Rehai with culture independent approach.A phylogenetic tree was constructed,among which two of these sequences are *Bacillus*,the other three sequences belong to Genera *Hydrogenobacter*,*Pseudomonas* and a potential genus that belongs to the family *Thermodesulfobacteriaceae*,it takes a position between the Genera *Geothermobacterium* and *Thermodesulfobacteria*.

Key words: hot spring culture-independent approach 16 S rDNA clone

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