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论著

山丘疫区杜氏利什曼原虫核糖体基因内转录间隔区的克隆及序列分析

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摘要

目的 构建我国山丘疫区杜氏利什曼原虫分离株前鞭体核糖体DNA(rDNA)内转录间隔区(ITS)片段克隆,并进行测序及同源性分析。 方法 提取杜氏利什曼原虫前鞭毛体DNA进行PCR扩增,将扩增出rDNAITS片段克隆入pMD18 Tvector上,双脱氧链末端终止法测序。 结果 扩增出约 1000bp的rDNAITS片段。测序结果表明山丘疫区的2株利什曼原虫L.d.SC10和L.d.6分别为1027bp和1028bp。序列分析结果表明,L.d.SC10和L.d .6有一定差异。 结论 获得了我国山丘疫区杜氏利什曼原虫分离株L.d.SC10和L.d.6的前鞭体rDNAITS序列。

关键词 利什曼原虫 内转录间隔区 基因克隆 序列分析

分类号

Cloning and Sequence Analysis of the Ribosomal DNA ITS Gene of Leishmania donovani Isolates from Hill Foci of China

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Abstract

Objective To determine the nucleotide sequence of the ITS (internal transcribed spacer) gene of Leishmania donovani isolates from hill foci (L.d.SC10 and L.d.6), and to find out the difference of the gene sequences between the two isolates. Methods Specific ITS fragments from nuclear DNA of two Leishmania isolates were amplified by PCR, cloned into pMD18 T vector, and finally sequenced by the dideoxy chain termination method. Results Sequence analysis showed that the amplified DNA fragments of the two isolates were 1 027 bp (L.d.SC10) and 1 028 bp (L.d.6) respectively, showing a sequence difference. Conclusion Sequence difference exists between the Leishmania isolates L.d.SC10 and L.d.6 from hill foci in China.

Key words *Leishmania donovani* ITS Gene cloning Sequence analysis

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