

论著

福建棘隙吸虫与相关虫种随机引物扩增多态DNA分析及感染实验观察

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摘要

目的:探讨福建棘隙吸虫与相关虫种分类问题。方法:应用28个随机引物扩增多态性DNA,根据福建棘隙吸虫、貌小棘隙吸虫(安徽)与日本棘隙吸虫(福建和江西)作相似性分析;现场采检叶形棘隙吸虫(湖北)、貌小棘隙吸虫(安徽)和福建棘隙吸虫囊蚴进行人工感染实验及形态观察。结果:分析福建、安徽与江西省3种棘隙吸虫的469个多态DNA片段,福建棘隙吸虫与日本棘隙吸虫福建株、日本棘隙吸虫江西株与福建株、福建棘隙吸虫与安徽的貌小棘隙吸虫间基因组DNA的RAPD相似率分别为20.8%、99.7%及97.6%。湖北、安徽和福建3省均分别获得叶形棘隙吸虫、福建棘隙吸虫及日本棘隙吸虫3种棘隙吸虫。结论:福建棘隙吸虫与日本棘隙吸虫为两个独立虫种,安徽的貌小棘隙吸虫与福建棘隙吸虫为同一虫种。湖北、安徽与福建3省均存在叶形棘隙吸虫、福建棘隙吸虫和日本棘隙吸虫3种吸虫的混合感染,其中福建棘隙吸虫较为常见。

关键词 [棘隙吸虫](#) [随机扩增多态DNA](#) [感染实验](#) [虫种](#)

分类号

TAXONOMIC STUDIES ON ECHINOCHASMUS FUJIENENSIS AND ITS RELATED SPECIES BY RANDOM AMPLIFIED POLYMORPHIC DNA ANALYSIS AND EXPERIMENTAL INFECTION

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Abstract

AIM: To explore the identification and differentiation of *Echinochasmus fujianensis*, *Echinochasmus japonicus*(Jiangxi strain and Fujian strain), *Echinochasmus liliputanus* Anhui and *Echinochasmus perfoliatus* Hubei. METHODS: Random amplified polymorphic DNA analysis (RAPD) and experimental animal infection were performed. RESULTS: 469 polymorphic DNA fragments were obtained by 28 primers from 4 *Echinochasmus* species and strains in Fujian, Anhui and Jiangxi. 20.8% and 97.6% of the fragments in *Echinochasmus fujianensis* were the same as those in *Echinochasmus japonicus* Fujian strain and in *Echinochasmus liliputanus* Anhui, respectively. 99.7% of the fragments were the same between *Echinochasmus japonicus* Jiangxi strain and Fujian strain. CONCLUSION: *Echinochasmus fujianensis* and *Echinochasmus liliputanus* Anhui are the same species. *Echinochasmus fujianensis* is an independent species different from *Echinochasmus japonicus*. Polyinfection of *Echinochasmus fujianensis*, *Echinochasmus japonicus* and *Echinochasmus perfoliatus* exist in all the 3 provinces, Hubei, Anhui and Fujian, of which *Echinochasmus fujianensis* is a dominant species.

Key words [Echinochasmus fujianensis](#) [random amplified polymorphic DNA](#) [experimental infections](#) [species](#)

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