研究报告

基于线粒体控制区的滇金丝猴群体遗传学研究

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收稿日期 2005-4-12 修回日期 2006-2-12 网络版发布日期 2006-4-7 接受日期

摘要

滇金丝猴(Rhi nopi thecus bi eti)是我国著名的濒危保护动物。迄今为止,关于滇金丝猴并基于DNA序列的的群 体遗传学研究还没有报道。 本文测定了来自于云南省维西县滇金丝猴群体样本的线粒体控制区全序列以及部分个 体的细胞色素b全序列。在排除了核线粒体假基因存在的可能性之后,滇金丝猴维西群体内部被确认存在着两个序 ▶ 复制索引 列分歧较大的分枝。 即使如此,如果考虑到群体结构和迁移的影响,维西群体的遗传多样性水平可能并不高。

关键词 滇金丝猴 线粒体控制区 遗传多样性 群体结构 分类号 0953

Population Genetics of *Rhinopithecus bieti*: a Study of the Mitochondrial **Control Region**

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Abstract

Yunnan snub-nose monkey (<1>Rhinopithecus bieti</1>) is a famous endangered primate in China. So far, however, studies on its population genetics based on DNA sequences are not available. In this paper, the whole mitochondria control region of the samples from Weixi, Yunan Province as well as the whole cytochrome b gene in some individuals were sequenced. A deep divergence was observed within the Weixi population, which was confirmed after excluding the possibility of it being a nuclear pseudogene. Nonetheless, if the effects of population structure and migration are considered, the true level of polymorphism of the Weixi population may be not as high as observed.

Key words Yunnan snub-nose monkey mitochondria control region genetic polymorphism population structure

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