论著

子一代实验室钉螺细胞色素c氧化酶I、细胞色素b基因序列分析

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目的 了解钉螺子一代(F1)实验室钉螺群(湖北钉螺湖北亚种)内的遗传变异。 方法 实验室饲养的 F1钉螺,单个抽提基因组DNA,PCR扩增细胞色素c氧化酶I(COI)、细胞色素b(Cytb)基因并测序,用 GENETYX MACver.9软件进行同源排序、DNA和氨基酸序列分析,同时与GenBank相同基因序列比较。 结果 实验室螺群内,COI基因差异为12.2%,94个氨基酸发生变化。Cytb基因差异6.4%,有25个氨基酸发生变化。湖北亚种与滇川亚种COI基因序列差异率为 13.5%。与GenBank中湖北钉螺滇川亚种 Cytb基因序列比较,差异为 13.6%,在 203个氨基酸中 6个氨基酸发生变化。 结论 F1实验室钉螺群内核苷酸与氨基酸序列均发生变异。

关键词 螺 子代 基因 遗传变异

分类号

Analysis of Cytochrome c Oxidase I and Cytochrome b Genes of F1 in Laboratory Line Oncomelania hupensis hupensis

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Abstract

Objective To analyze the diversity of F1 in laboratory line Oncomelania hupensis hupensis. Methods Genomic DNA was isolated, and cytochrome c oxidase I (COI) and cytochrome b (Cytb) genes were amplified by polymerase chain reaction (PCR). The nucleotide sequences of the PCR products were analyzed by GENETYX MAC software package (ver.9). Results Pairwise divergences among six F1 individuals were found in 12.2% nucleotides in COI gene fragments. 13.5% of genetic divergences between O.h.hupensis and O.h.robertsoni were identified. 94 amino acids were observed in difference. In Cyt b gene fragments, 6.4% of divergence was observed among four F1 samples. 25 amino acids were observed differently. The divergence of Cyt b genes from O.h.robertsoni and O.h.hupensis were 13.6%, included 6 amino acids. Conclusion Diversities were found in both COI and Cyt b gene sequences of F1 in laboratory line O.h.hupensis.

Key words Snails F1 filial generation Genes Genetic variation

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