水松自然种群和人工种群遗传多样性比较

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Genetic diversity of natural and planted Glyptostrobus pensilis populations: A comparative study.

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摘要

采用ISSR分子标记技术分析水松不同起源种群的遗传多样性.结果表明:10条引物共检测出95个扩增位点,多态位点数占39.0%.与 其他濒危裸子植物相比,水松的遗传多样性较低,遗传分化系数G_{st}为0.3982,基因流N_m仅0.3778,种群间存在一定程度的遗传分化, 但种群内变异占主导地位;遗传距离与地理距离呈正相关关系,自然种群的多态位点百分率(P)、Nei的条带多样度(Ha)和Shannon 信息指数(/)平均值(39.3%、0.1499和0.2202)分别高于人工种群(30.7%、0.1265和0.1759).自然种群的遗传分化系数 (G_{st} =0.4513) 和平均遗传距离(D=0.0301)也高于人工种群(G_{st} =0.3025,D=0.0192).

关键词: 水松 遗传多样性 遗传结构 自然种群 人工种群 ISSR

Abstract:

Glyptostrobus pensilis is a rare and endangered relict species in China. To make a comparative study on the genetic diversity and genetic structure of

natural and planted G. pensilis populations would have significance in the conservation and proliferation of the species. Samples from the main distribution regions of G. pensilis were analyzed by ISSR (inter simple sequence repeats) molecular marker. A total of 95 discernible DNA fragments were detected with 10 ISSR primers, of which, polymorphic loci occupied 39.0%, suggesting that the genetic variation in the test G. pensilis populations was at a very low level, compared with other endangered gymnosperm. The genetic differential index $(G_{\rm st}=0.3982)$ and the gene flow $(N_{\rm m}=0.3778)$ indicated that there existed genetic differentiation among populations but the differentiation dominated within populations. There was a significant positive correlation between genetic distance and geographical distance. The mean values of polymorphic loci (P), Nei's gene index (H_p) , and Shannon information index (I) of natural populations (P=39.9%, H_p =0.1499, I=0.2202) were much higher than those of planted G. pensilis populations (P=30.7%, H_p =0.1265, I=0.1759), and the coefficient of gene differentiation (G_{st}) and genetic distance (D) of natural populations (G_{st} =0.4513, D=0.0301) were also much higher than those of the planted populations (G_{st} =0.3025, D=0.0192).

Key words: Glyptostrobus pensilis genetic diversity genetic structure natural population planted population **ISSR**

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