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一种改良的水稻细胞质基因组制备方法

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An Improved Method for Extraction of Rice Cytoplasmic Genome

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摘要

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摘要 水稻叶绿体和线粒体基因组较小,且均被测序清楚,可以作为研究细胞质遗传的良好材料,而如何快速有效地分离纯化获得高产量和高质量的细胞质基因组是从DNA水平上研究细胞质遗传变异的前提条件。本研究结合了蔗糖密度梯度离心法-全基因组DNA扩增法(whole genome amplification, WGA),对细胞质基因组的制备进行了改良。改良后的方法仅使用 20 g 的水稻叶片,即可在 2 d 时间内得到浓度达 300 ng/μL 以上,总量达 40 μg 以上的高纯度(OD 260/280 值为1.8~2.0)、完整的叶绿体DNA (chloroplast DNA, cpDNA)和线粒体DNA (mitochondrial DNA, mtDNA)。该法制备的细胞质基因组可以满足多种细胞质基因组实验的要求,包括Solexa全基因组测序技术的要求。

关键词: 水稻 叶绿体DNA 线粒体DNA

Abstract: Due to the small size of rice chloroplast and mitochondria genomes, they are good models for the study on extranuclear inheritance. The key step is how to separate and purify the highquality cytoplasmic genome. In this research, we have developed a new technique based on combination of sucrose stepgradient centrifugation and whole genome amplification (WGA). Through the developed method, we can get high quality and quantity (OD 260/280=18~20) DNA and mitochondrial DNA with about 300 ng/μL of concentration and 40μgof grosses from only 20grice leaves comparing with traditional methods. The chloroplast DNA and mitochondrial DNA can fulfill the needs of much cytoplasmic genome study, even the needs of Solexa whole genome sequencing (new generation sequencing technology).

Keywords: rice chloroplast DNA mitochondrial DNA

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