

菠菜种质遗传多样性和亲缘关系的AFLP 分析

吴娅妮, 康俊根, 王文科, 孟淑春

(1 北京市农林科学院蔬菜研究中心, 农业部华北地区园艺作物生物学与种质创制重点实验室, 农业部都市农业(北方)重点实验室, 北京 100097; 2 山西师范大学生命科学学院, 山西临汾 041004)

Genetic Diversity and Relationship of Spinach Germplasm Revealed by AFLPs

WU Ya-Ni, KANG Jun-Gen, WANG Wen-Ke, MENG Shu-Chun

(1Beijing Vegetable Research Center, Beijing Academy of Agriculture and Forestry Sciences, Key Laboratory of Biology and Genetic Improvement of Horticultural Crops (North China), Ministry of Agriculture, Key Laboratory of Urban Agriculture (North), Ministry of Agriculture, Beijing 100097, China; 2College of Life Science, Shanxi Normal University, Linfen, Shanxi 041004, China)

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摘要 利用AFLP 标记技术对110 份来源不同的菠菜 (*Spinacia oleracea* L.) 种质进行遗传多样性分析, 结果表明, 筛选出的20 对EcoR I /Mse I 引物组合共扩增出882 条带, 其中208 条多态性条带, 多态性比率23.6%; 种质间的遗传相似系数为0.64 ~ 0.87, 不同来源组群的Nei' s 遗传多样性指数范围为0.1887 ~ 0.2501, 总体为0.2830, Shannon 信息指数范围为0.2789 ~ 0.3793, 总体为0.4337. 种质间

基于遗传相似系数的UPGMA 聚类分析、主坐标分析与组群间基于Nei' s 遗传距离的聚类分析结果基本相同, 与地理来源有很高的一致性。全部供试种质可分为两类, 欧美、西亚、东亚及中国北方种质聚为一类, 部分日本种质和中国的南方种质聚为另一类, AFLP 标记能很好地从分子水平揭示菠菜资源的亲缘关系。由亲缘关系推测中国的南、北方菠菜种质可能有着不同的起源。

关键词: 菠菜 遗传多样性 亲缘关系 AFLP

Abstract: Amplified fragment length polymorphism (AFLP) markers were employed to assess the genetic diversity and relationship of 110 spinach (*Spinacia oleracea* L.) germplasms from different geographical origins. Among the 882 polymorphic bands obtained from 20 selective primer pairs, 228 (23.6%) of them was polymorphic. The genetic similarity ranged from 0.64 to 0.87 for all germplasms accessions. Nei' s gene diversity index ranged from 0.1887 to 0.2501 for different populations, with the total value of 0.2830. Shannon information index varied from 0.2789 to 0.3793, with the total value of 0.4337. UPGMA Cluster suggested genetic relationship of all the accessions were significantly related to their geographical origins. The results of PCA, UPGMA and based Nei' s gene diversity of populations clustering analysis exhibited a good consistency with UPGMA, and were also consistent with geographical origins. All the accessions were classified and clustered into two major groups. Group 1 was comprised of the spinach originated in European, America, West Asia, East Asia and Northern China, whereas group 2 was comprised of the spinach originated in the Southern China and Japan. The results suggested that Northern and Southern Chinese spinach population may have different origins.

Keywords: *Spinacia oleracea* L., genetic diversity, genetic relationship, AFLP

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