

Abstract

Date Palm (*Phoenix dactulifera* L.) is monocotyledone, dioecious and long life which could be highly economically important. Assigning the granule giver and the genetic distance between the male and female cultivars are the most important goal of this research. This research was carried out at the Biotechnology laboratory of agriculture research center in Hormozgan. Province in order to do analysis the genetic diversity of 14 male and 26 female cultivars of Date palm by using the indicators of microsatellite. DNA extracted of young leaves via Sambrook CTAB procedure (1998) with little modification and DNA amplification was carried out using 10 microsatellite primer pairs. PCR products were separated on a 8% undenaturated acrylamide gel containing 7M Urea and were stained by silver staining method. Gels were then scored based on either the presence or absence of the bands. Polymorphism information content (PIC) for each SSR marker was determined. PIC was 0.63 for HQ542225 primer to 0.85 for HQ542208 primer with a mean value of 0.77. Number of alleles varied between 5-9 and total of 76 alleles were identified with an average of 7.6 alleles per locus. Genetic relationships among cultivars were represented by a dendrogram based on the Nei's Genetic similarity coefficient and Neighbor-joining method which was considered for cluster analysis. Cluster analysis divided the 40 Date palm cultivars in three major groups. The cultivars were categorized and classified into different groups by using geographical distribution. The highest similarity coefficient value was observed between Khanizi and Almehtary. The minimum of similarity was obtained between Abonarenja and other cultivars. Results show that SSR markers are useful tools for genetic diversity fingerprinting of Date palm.

Key words: SSR markers, male and female cultivars, date palm, genetic diversity



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Investigation of genetic diversity of date palm cultivars in Sistan and Baluchistan and Hormozgan provinces using SSR markers

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