

## MOLECULAR MARKERS AS A BOON FOR PLANTS GENETIC FIDELITY: A REVIEW

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### ABSTRACT

Molecular markers have brought to the fore an extremely efficient method of estimating the genetic diversity of germplasm collections of various plant species. They can enhance the speed and effectiveness of plant breeding, and can even be used efficiently for quality control. Different types of molecular markers have made it possible to detect the sequence of DNA or protein whose inheritance can be monitored and thus proves an important tool for studying genetic fidelity among plants. These markers are extremely essential for phylogenetic analysis adding new dimension to the evolutionary theories. The present review deals with the different types of molecular markers and their possible implications in improving the breeding process, for sex identification of dioecious plants and testing the genetic fidelity in plants.

**Keywords :** molecular markers, plant breeding, genetic fidelity, germplasm.

### REFERENCE

- Andersen J R, and Lubberstedt T. 2003. Functional markers in plants. Trends in Pl.Sc. 8(11): 554-560
- Bernatzkry., And S. D. Tanksley, 1986 Toward a saturated linkage map of tomato based on isozymes and random cDNA sequences. Genetics 112: 887-898.
- Beckmanjn., S., and Soller M. 1986 Restriction fragment length polymorphisms in plant genetic improvement. Oxf. Surv. Plant Mol. Cell Biol. 3: 197-250.
- Crowe M.L., Serizet,C., Thareau,V., Aubourg,S., Rouzé,P., Hilson,P., Beynon,J., Weisbeek,P., van Hummelen,P., Reymond,P. *et al.* (2003) CATMA—A complete *Arabidopsis* GST database. *Nucleic Acids Res.*, 31, 156–158.
- Daniel T. Busemeyer, Stephan Pelikan, Robert S. Kennedy and Steven H. Rogstad 1997. Genetic Diversity of Philippine *Rubus moluccanus* L. (Rosaceae) Populations Examined with VNTR DNA Probes. *Journal of Tropical Ecology*, Vol. 13, No. 6 :867-884
- Egea-Gilabert C, M.J. Dickinson, G. Bilotti and M.E. Candela 2003. Isolation of Resistance Gene Analogs in Pepper Using Modified AFLPs. *Biologia Plantarum* 47 (1):27-32.
- Fredslund J, Madsen LH, Hougaard BK, Sandal N, Stougaard J, Bertoli D, Schauer L (2006) GeMprospector - online design of cross-species genetic marker candidates in legumes and grasses. *Nucleic Acids Res* 34: W670–W675

- Fjellheim S, Rognli OA (2005). Genetic diversity within and among Nordic meadow fescue (*Festuca pratensis* Huds.) cultivars determined on the basis of AFLP markers. *Crop Sci.* 45 (5): 2081-2086.
- Gale, M. D. and Devos, K. M. 1998. Comparative genetics in the grasses *Proc. Natl. Acad. Sci. USA*, 1971–1974.
- He JP.2009. Amplified consensus genetic markers and its application in plants. *Yi Chuan.*;31(9):913-20. Review. Chinese.
- Hei.Enl:Jaris, T., Slocums M., . Wright,A . Schaefer and J. Nienhuis, 1986 Construction of genetic linkage maps in maize and tomato using restriction fragment length polymorphisms. *Theor. Appl. Genet.* 72: 761-769.
- James KE, Schneider H, Ansell SW, Evers M, Robba L, et al. 2008 Diversity Arrays Technology (DArT) for Pan-Genomic Evolutionary Studies of Non-Model Organisms. *PLoS ONE* 3(2): e1682. doi:10.1371/journal.pone.0001682
- Jonathan H. Crouch Hutokshi K. Crouch Abdou Tenkouano Rodomiro Ortiz 1999. VNTR-based diversity analysis of 2x and 4x full-sib *Musa* hybrids. *Electronic Journal of Biotechnology* [online]. Vol.2 No.3, Issue of December 15, 1999. Available from: <http://www.ejbiotechnology.info/content/vol2/issue3/full/3/index.html>. ISSN 0717-3458.
- Kwok PY, Chen X. Detection of single nucleotide polymorphisms. *Curr Issues Mol Biol.* 2003;5:43–60.
- Rostoks N, Borevitz JO, Hedley PE, Russell J, Mudie S, Morris J, Cardle L, Marshall DF and Waugh R 2005. Single-feature polymorphism discovery in the barley transcriptome. *Genome Biol.* 2005; 6(6): R54.
- Steven H. Rogstad, Brian Keane and John Beresh 2002. Genetic Variation across VNTR Loci in Central North American *Taraxacum* Surveyed at Different Spatial Scales. *Plant Ecology*, Vol. 161(1):111-121 Wenzl P, Carling J, Kudrna D, et al. (June 2004). “Diversity Arrays Technology (DArT) for whole-genome profiling of barley”. *Proc. Natl. Acad. Sci. U.S.A.* 101 (26): 9915–20.
- Tanksley S. D, Young N. D., Paterson A. H. & Bonierbale M. W. 1989. RFLP Mapping in Plant Breeding: New Tools for an Old Science. *Nature Biotechnology* 7: 257 – 264.
- Theresa M. Fulton, Rutger Van der Hoeven, Nancy T. Eannetta and Steven D. Tanksley 2002. Identification, Analysis, and Utilization of Conserved Ortholog Set Markers for Comparative Genomics in Higher Plants. *The Plant Cell*, Vol. 14, 1457-1467
- Thareau V, Déhais P., Serizet C., Hilson P., Rouzé P. and Aubourg S. (2003) Automatic design of gene-specific sequence tags for genome-wide functional studies. *Bioinformatics*, 19: 2191-2198.
- Varshney RK, Graner A, Sorrells ME 2005. Genic microsatellite markers in plants: features and applications. *Trends Biotechnol.* 23(1):48-55
- Vos P, Hogers R, Blecker M, Rijans M, van der Lee T, Hornes M, Frijters A, Pot J, Poleman J, Kuiper M, Zabeau M (1995) AFLP: a new technique for DNA fingerprinting. *Nucleic Acids Res* 23:4407–4414

- Weeden N F, Hemmatt M, Lawson D M, Lodhi M, Bell R L, Manganaris A G, Reischs BI, Brown SK and Ye.GN 1994. Development and application of molecular marker linkage maps in woody fruit crops. *Euphytica* 77 (1-2)
- Wenzl P, Jason Carling J, Kudrna D, Jaccoud D, Huttner E, Kleinhofs A, and Kilian A 2004. Diversity Arrays Technology (DArT) for whole-genome profiling of barley PINAS-101(26): 9915-9920
- Yong-Quan LU, Zi-Hong YE and Wei-Ren WU 2006. Analysis of the Phylogenetic Relationships Among Several Species of *Gramineae* Using ACGM Markers. *Acta Genetica Sinica* 33 (12): 1127-1131
- Yoshimura A, Ideta O, Iwata N 1997. Linkage map of phenotype and RFLP markers in rice. *Plant Mol Biol.*;35(1-2):49-60.
- Zhou Z., Bebeli P. J., Somers D. J. and Gustafson J. P. 1997. Direct amplification of minisatellite-region DNA with VNTR core sequences in the genus *Oryza* . *Theor Appl Genet* 95 : 942-949