

# Plant Breeding: A Structural Frame Work In Agriculture

Nischita. P

Department of Biotechnology, Centre for Post Graduate  
Studies, Jain University, Jayanagar,  
Bengaluru-560011, India

M. R. Dinesh

Division of Fruit Crops, ICAR-Indian Institute of  
Horticultural Research, Hessaraghatta Lake Post,  
Bengaluru-560089, India

**Abstract:** *Breeding in plants is recognized as one of the oldest tool of agricultural development initiated by humans over decades. The evolving of civilization manifested the achievement in plant breeding though not been acknowledged by common man, the reason being poor understanding of breeding concept. In spite of good outcome from traditional methods; new technology with novel inventions are emerging consistently and furthermore advance research is necessary to fulfill the sustainability of human, thus producing improved cultivars by identifying the desirable ones with combination of characters for better expression pattern. This review talks about the principles, methodology, current approach and future outlook of plant breeding, more importantly in crop improvement.*

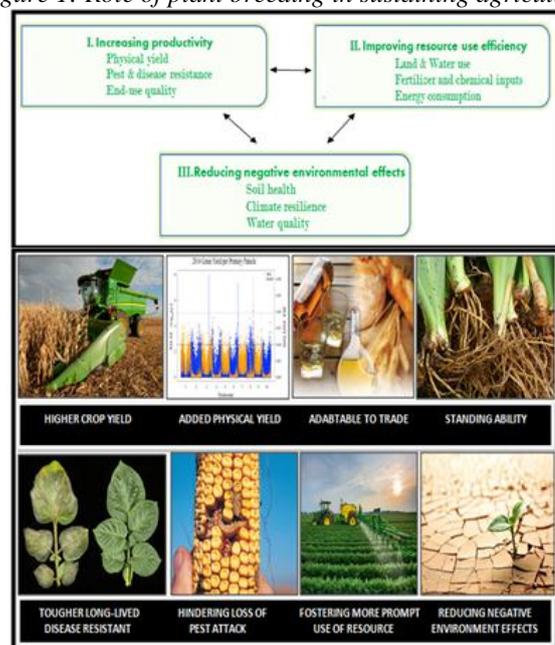
**Keywords:** *Breeding, Agriculture, Genetics, Crop Improvement, DNA Technology*

## I. INTRODUCTION

Although the requirement of food is been rapidly increasing due to flourishing human population there needs to be an improvement in quality of food especially with enhanced nutritive value along with decreased farming inputs thereby avoiding the environment pollution especially triggered by emissions of CO<sub>2</sub> and nitrogenous fertilizers (Mark and Peter, 2000). The challenging efforts are being made these days to increase yield and resisting environmental threats with global change. This change in food security calls for exploitation of revised technologies to secure food supply, thus considering various criteria namely crop yield and expenditures (Goldman, 2013). Several molecular genetics parameters, recent advancement in integrative genomics, bioinformatics have into the domain of molecular breeding is been formulated (Ghanemi, 2013a; Young, 1999; Cardon and Palmer, 2003; Ramanna and Jacobsen, 2003; Chen et al., 2013; Whitford et al., 2013, Li et al., 2012). The plant breeding industry is a major contributor to more sustainable agriculture and food production (Figure 1). This review furnishes and highlights the current approaches along with future prospects in some of the instructive model leading to crop improvement and environmental stress resistance. Additionally plant breeding coupled with extensive genomic

resources by understanding the concept of emerging model plant thus providing new opportunities, innovative approaches to meet challenges ahead.

Figure 1: Role of plant breeding in sustaining agriculture





- [5] Cardon, L. R., and Palmer, L.J, (2003) .Population stratification and spurious allelic association. *Lancet* 361(9357).; 598-604.
- [6] Ramanna, M.S and Jacobsen, E. (2003) . Relevance of sexual polyploidization for crop improvement – A review. *Euphytica*.133(1): 3-8.
- [7] Chen, J, Xue, B. Xia., and X. Yin W. (2013). A novel calcium-dependent protein kinase gene from *Populus euphratica*, confers both drought and cold stress tolerance. *Biochem Biophys Res Commun*. 22;441(3):630-6. doi: 10.1016/j.bbrc.2013.10.103.
- [8] Whitford, R. Fleury, D. Reif. J. C. Garcia, M. Okada, T. Korzun, V., and Langridge., P .Hybrid breeding in wheat: technologies to improve hybrid wheat seed production. *J Exp Bot*. 2013; 64(18):5411-28. doi: 10.1093/jxb/ert333.
- [9] Li, X. Zhu, C. Wang, J., and Yu, J. (2012)Chapter six - Computer Simulation in Plant Breeding. *Advances in Agronomy*. L. S. Donald, Academic Press.116: 219-264.
- [10] King, Z. R. Bray, A. L. Lafayette, P. R., and Parrott, W. A. (2013)..Biolistic transformation of elite genotypes of switchgrass (*Panicum virgatum* L.). *Plant Cell Rep*.33(2):313-22, doi: 10.1007/s00299-013-1531-1.
- [11] Alheit, K. V. L. Busemeyer, W. Liu, H .P. Maurer, Gowda, V. Hahn, S. Weissmann, A .Ruckelshausen, J. C. Reif., and Wurschum, T . Multiple-line cross QTL mapping for biomass yield and plant height in triticale (x *Triticosecale* Wittmack). *Theor Appl Genet* 2013; 127(1) pp 251–260.
- [12] Chao, M. Yin, Z. Hao, D. Zhang, J. Song, H. Ning, A. Xu, X., and Yu, D. (2013). Variation in Rubisco activase (RCA $\beta$ ) gene promoters and expression in soybean *Glycine max* (L.) Merr.]. *J Exp Bot* 65:47–59. doi:10.1093/jxb/ert346.
- [13] Alessandro, T. X. X. Marc, M. Rajiv, S. Florian, S. Christina, Ingvarlsen, O. M. Jordi, C. Joanne, R. Robbie, W. Alan, H.S. Klaus Pillen, S. K .R. Benjamin, K. Luigi. C. William, T. B. Thomas., and Flavell., A J.(2013) Structural and temporal variation in genetic diversity of European spring 2-row barley cultivars and association mapping of quantitative traits. *The Plant Genome*. doi: 10.3835/plantgenome2013.03.0007.
- [14] Zhang, K. J. Wang, L. Zhang, C. Rong, F. Zhao, T. Peng, H. Li, D. Cheng, X. Liu, H. Qin, A. Zhang, Y. Tong., and Wang, D.(2013) Association analysis of genomic loci important for grain weight control in elite common wheat varieties cultivated with variable water and fertiliser supply. *PLoS One*. 8(3): e57853.
- [15] Hull, R .Chapter 15 - Plant Viruses and Technology. *Plant Virology* (Fifth Edition). R. Hull. Boston, Academic Press: 2014;877-926.
- [16] Ghanemi, A. (2013b). Psychiatric neural networks and neuropharmacology: Selected advances and novel implications. *Saudi Pharmaceutical Journal* <http://dx.doi.org/10.1016/j.jsps.2013.01.008>.
- [17] Boubertakh ,B, Liu, X. G, Cheng, X. L., and Li P.(2013) A Spotlight on Chemical Constituents and Pharmacological Activities of *Nigella glandulifera* Freyn et Sint Seeds. *Journal of Chemistry*: 12; <http://dx.doi.org/10.1155/2013/820183>.
- [18] Cheng, X. L. Qi, L.W. Wang, Q. Liu, X. G. Boubertakh, B. Wan, J. Y. Liu, E. H., and Li, P. (2013). Highly efficient sample preparation and quantification of constituents from traditional Chinese herbal medicines using matrix solid-phase dispersion extraction and UPLC-MS/MS." *Analyst*.138(8): 2279-2288.
- [19] Lee, C. J. Wu, Y. T. Hsueh, T. Y. Lin, L. C. and Tsai, T. H. (2013). Pharmacokinetics and oral; 28(5):630-6. doi: 10.1002/bmc.3081.