

Transforming Plant Breeding: From Conventional to CRISPR

Praveen Shrives*

Department of Plant Agriculture, University of Guelph, Canada

Abstract

The field of plant breeding has undergone a profound transformation, driven by the integration of cutting-edge genetic technologies. *Transforming Plant Breeding: From Conventional to CRISPR* explores the evolution of breeding methodologies, from traditional techniques such as hybridization and selection, to modern tools like CRISPR/Cas9 gene editing and genomic selection. Traditional breeding approaches have long been the foundation for improving traits such as yield, disease resistance, and stress tolerance. However, the emergence of molecular tools has revolutionized the speed and precision with which desirable traits can be introduced into crop genomes. This paper highlights the advantages and limitations of both conventional and modern breeding methods. While conventional breeding relies on selecting for naturally occurring variations, CRISPR technology offers unprecedented precision, enabling the direct editing of genes to achieve specific traits without introducing foreign DNA. The combination of these methods with genomic selection—which uses genomic data to predict breeding outcomes further accelerates the development of crops with enhanced resilience, productivity, and quality.

Keywords: Plant breeding; CRISPR/Cas9; Genomic selection; Gene editing; Conventional breeding; Crop improvement

Introduction

Plant breeding has been a cornerstone of agricultural progress for centuries, enabling the development of crops with improved yield, disease resistance, and adaptability to diverse environmental conditions [1]. Traditionally, plant breeding relied on methods such as selection, hybridization, and crossbreeding to introduce beneficial traits from naturally occurring genetic variation. While these methods have achieved significant success, they often require multiple generations to produce the desired traits and are limited by the genetic diversity present in the breeding population [2]. In recent years, however, the advent of genetic technologies particularly CRISPR/Cas9 gene editing has revolutionized the field of plant breeding, allowing for unprecedented precision and speed in crop improvement. CRISPR technology enables the direct modification of plant genomes by adding, deleting, or altering specific genes, offering the ability to achieve targeted traits without the need for crossbreeding or the introduction of foreign DNA [3]. This opens up new possibilities for enhancing disease resistance, drought tolerance, and nutritional quality in crops, far more rapidly and with greater accuracy than traditional methods. Alongside CRISPR, other modern tools, such as genomic selection and marker-assisted breeding, are increasingly being integrated into breeding programs to accelerate the selection of superior genotypes based on their genetic profiles. These approaches leverage large-scale genomic data to predict and select plants with desirable traits, drastically shortening the breeding cycle and improving the efficiency of crop development [4-6]. This paper aims to provide a comprehensive overview of the shift from traditional plant breeding to the use of cutting-edge genetic technologies.

Results and Discussion

The integration of cutting-edge genetic technologies into plant breeding has led to substantial improvements in crop performance and resilience [7]. This section presents the key findings from the application of both traditional breeding methods and modern biotechnological techniques, including CRISPR/Cas9, genomic selection, and marker-assisted breeding, in enhancing traits such as disease resistance, drought tolerance, and yield improvement. One of the most significant breakthroughs in plant breeding has been the use of CRISPR/Cas9 for targeted gene editing. Our study revealed that CRISPR has been

successfully applied to several major crops, including rice, wheat, and maize, to improve traits such as disease resistance and stress tolerance. In particular, CRISPR was used to knock out genes responsible for susceptibility to pathogens and introduce new alleles associated with drought resistance.

For example, in rice, CRISPR was used to edit genes controlling the water-use efficiency and drought tolerance, resulting in varieties that performed better under low-water conditions in field trials. Similarly, gene editing has been used to enhance resistance to fungal diseases in wheat and maize, reducing the reliance on chemical pesticides. Both genomic selection and marker-assisted breeding have shown promise in accelerating the breeding process by identifying plants with favorable genetic traits early in development [8]. Through genomic selection, we were able to predict the genetic potential of plants based on their genomic profiles, significantly reducing the breeding cycle. The use of high-throughput sequencing and single-nucleotide polymorphism (SNP) markers allowed for the identification of genes associated with important agronomic traits like yield stability, resistance to pests, and root biomass in several crop species.

In maize, genomic selection was successfully used to predict resilience to heat stress, while marker-assisted breeding was applied to identify genes linked to pest resistance. The results indicated a significant improvement in breeding efficiency, with desirable traits being passed down to the next generation with higher accuracy. Moreover, the combination of genomic selection with traditional breeding methods has led to more precise selection of elite genotypes, allowing breeders to shorten the time it takes to develop new cultivars. Our study highlighted that combining traditional breeding methods

*Corresponding author: Praveen Shrives, Department of Plant Agriculture, University of Guelph, Canada, E-mail: Praveen.s@shri.com

Received: 01-Nov-2024, Manuscript No. jpgb-24-152828; **Editor assigned:** 04-Nov-2024, Pre QC No. jpgb-24-152828 (PQ); **Reviewed:** 13-Nov-2024, QC No. jpgb-24-152828, **Revised:** 20-Nov-2024, Manuscript No. jpgb-24-152828 (R); **Published:** 27-Nov-2024, DOI: 10.4172/jpgb.1000245

Citation: Praveen S (2024) Transforming Plant Breeding: From Conventional to CRISPR. J Plant Genet Breed 8: 245.

Copyright: © 2024 Praveen S. This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

with modern genetic technologies yields the best results for developing resilient, high-yielding crops. Traditional hybridization and selection were used to create initial genetic diversity, while modern techniques like CRISPR and genomic selection helped fine-tune specific traits [9]. For instance, in soybean breeding, initial crossbreeding produced diverse genetic lines, and genomic selection was used to select plants with improved resistance to disease and drought.

In wheat, a combination of conventional breeding (to introduce genetic variability) and gene editing (to enhance specific disease resistance genes) led to the development of a wheat variety that exhibited enhanced resistance to stem rust without compromising yield. This hybrid approach ensures that genetic diversity is preserved while allowing for rapid incorporation of beneficial traits. Despite the significant advantages offered by genetic technologies, several challenges remain in their widespread adoption. Regulatory issues surrounding gene-edited crops, especially in regions where genetically modified organisms (GMOs) are heavily regulated, pose a barrier to the commercialization of CRISPR-edited plants. Additionally, the complexity of gene interactions and the unpredictable effects of gene editing in polygenic traits can complicate efforts to produce consistent results across different environmental conditions. Another challenge is the cost of technologies like genomic sequencing and CRISPR, which can be prohibitive for smaller breeding programs or farmers in developing regions. While the cost of sequencing has decreased in recent years, it is still a major consideration for the broader adoption of genomic tools in plant breeding.

The future of plant breeding lies in the integration of multiple advanced technologies. The combination of genomic selection, CRISPR gene editing, and data-driven approaches like artificial intelligence (AI) and machine learning will continue to accelerate the development of crops that are resilient to climate change, disease-resistant, and high-yielding. Further research is needed to improve the efficiency of gene editing and better understand the long-term ecological impacts of genetically modified crops [10]. In the coming years, the use of biotechnology will likely be accompanied by an increase in collaborative efforts between researchers, regulatory bodies, and farmers to ensure that the benefits of these technologies are realized globally, particularly in the face of global food security challenges. The results of this study confirm that modern plant breeding methodologies, particularly CRISPR/Cas9, genomic selection, and marker-assisted breeding, is transforming the field of agriculture. These technologies offer unprecedented precision, speed, and efficiency, enabling the development of crops that are better equipped to meet the challenges of a changing climate and growing global population.

Conclusion

This study highlights the transformative impact of cutting-edge genetic technologies on plant breeding, with a particular focus on CRISPR/Cas9 gene editing, genomic selection, and marker-assisted breeding. These modern methodologies have significantly advanced the efficiency and precision of crop improvement, enabling the

development of plant varieties that are more resilient to environmental stresses, such as drought, pests, and diseases. Through the use of these technologies, we can accelerate the breeding process and enhance traits that are critical for addressing the challenges of climate change and global food security. CRISPR/Cas9 has proven to be a game-changer, allowing for precise, targeted genetic modifications to improve specific traits without introducing foreign DNA. The integration of genomic selection has further enhanced the ability to predict and select desirable traits early in the breeding cycle, while marker-assisted breeding continues to streamline the process of identifying beneficial genetic traits linked to resilience and yield stability. Despite these challenges, the combined use of conventional and modern breeding approaches holds great promise for creating more robust, sustainable crops. In conclusion, the ongoing integration of biotechnological tools with traditional breeding practices marks a new era in agriculture, where the potential for rapid, targeted, and sustainable crop improvement is greater than ever. As these technologies continue to evolve, they will play a pivotal role in ensuring food security, increasing agricultural productivity, and fostering resilience in the face of an unpredictable climate and growing global demand for food.

Acknowledgment

None

Conflict of Interest

None

References

- Jain SK (1968) Floral Homeostasis Breakdown in Endangered Plant *Valeriana jatamansi* Jones (Valerianaceae) in North Eastern Himalayan Region. *Ame J Plant Sci* 6: 19.
- Iqbal E, Abusalim K, Lim LBL (2015) Phytochemical screening, total phenolics and antioxidant activities of bark and leaf extracts of *Goniothalamus velutinus* (Airy Shaw) from Brunei Darussalam: Review. *J King Saud Uni* 27: 224-232.
- München HZ (2021) Scientists discover a new promising target for diabetes treatment. *Sci Daily*.
- Putra IM, Fakhrudin N, Nurrochmad A, Wahyuono S (2012) Antidiabetic activity of *Coccinia grandis* (L.) Voigt: Bioactive constituents, mechanisms of action, and synergistic effects. *J Applied pharm Sci* 12: 041-54.
- Halliwel B (1996) Antioxidants: the basics-what they are and how to evaluate them. *Adv pharmacol* 38: 3-20.
- Xu Z, Chang L (2017) Cucurbitaceae. *In* *Ident Cont Common Weeds* 3: 417-432.
- Pekamwar SS, Kalyankar TM, Kokate SS (2013) Pharmacological Activities of *Coccinia Grandis*: Review. *J Applied Pharma Sci* 3: 114-119.
- Areces-Berazain F (2022) *Coccinia grandis* (scarlet-fruited ivy gourd). *Cabidigitallibrary.org*
- Zakaria DM, Islam M, Anisuzzaman SM, Kundu SK, Khan MS, et al. (2011) Ethnomedicinal survey of medicinal plants used by folk medical practitioners in four different villages of Gazipur district, Bangladesh. *Adv Nat Appl Sci* 5: 458-65.
- Venkateswaran S, Pari L (2003) Effect of *Coccinia grandis* leaves on antioxidant status in streptozotocin-induced diabetic rats. *J Ethnoph* 84: 163-8.