

Mechanisms and Role of Genetic Regulation in Improving Crop Yields

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Submitted: 2023, Oct 10; Accepted: 2023, Nov 08; Published: 2023, Dec 15

Citation: Kumsa, F., Kuma, T. (2023). Mechanisms and Role of Genetic Regulation in Improving Crop Yields. *J Gene Engg Bio Res*, 5(3), 245-250.

Abstract

As the population nowadays is booming increasing the amount of crop manufacturing would end in such lots of inconveniences. This can be obvious thanks to the actual fact the additional population will increase want, the greater yield might be accustomed satisfy food demand. To beat this kind of phenomenon, it miles too much encouraged to apply better yielded higher crops. However, most of the countries did no longer supply attention to growing and using genetically-regulated crops. This practice and therefore the use of genetically modified organisms are often hampered by limited resources and knowledge base. On the other hand, current temperature change, which threatens the globe, appears to pose a practical and potential danger to crop production, because it causes crop failures and associated yield declines. Current review papers are created from topic-related literature to produce summarized knowledge of the mechanisms of crop improvement and use.

Keywords: Crops, Demand, Gene Regulation, Population, Yield

1. Introduction

The world population is now growing much faster and faster than ever before in history. The results of growth could be a reduction in existing agricultural land and environmental degradation, which could lead to a general food deficit [1]. These common issues are addressed through a variety of mechanisms, including the use of plant varieties with high yield per unit area. The impact of genetic improvement on recent crop varieties production requires the identification of genetic diversity of morphological, structural, and physiological characteristics that affect yield. This is often associated with the manipulation of complex signs associated with plant growth and development [2,3]. Some characteristics of plant development, such as plant structure, leaf characteristics, and vascular structure, are the most important characteristics that determine crop yield (see Figure 1). The leaves are the site of photosynthesis and can be genetically engineered to improve photosynthetic yields. The plant's vasculature is another function that regulates the overall performance of the plant, providing not only mechanical strength, but also a conduit for the transport of water, minerals, and photosynthesis [4].

Regulation of gene expression is also recommended for yield production. This gene regulates cell number and organ size, which can increase yield. For example, isolated and defined, and determined

the gene (CNR1) from the maize cell count. Absolutely, CNR1 has been shown to reduce overall plant size when ectopically overexpressed, but the plant and organ size increased when its expression was co-suppressed or silenced. Genomic engineering may be a relatively new experience for improving crop yields, nutritional value, herbicide tolerance, biological and abiotic stress tolerance [5,6]. During this review, the mechanistic role of gene regulation in crops for better yields was evaluated.

1. Analysis of the Related Literature Reviews**1.1. Regulation and Expression of Plant Genes**

The genetic growth regulators of plants are present within the nucleus. The nucleus contains most of the functional genes of a plant in the genome, which is organized consistent with the final model of eukaryotes [7]. For several years, the regulation of organic phenomenon in plants has been studied at the phenotypic level. The biological revolution led to biochemical and molecular genetic techniques has displayed possibilities for understanding the links between genetics and physiology like genotype and phenotype. This includes observing how plants respond to external factors. However, a few years ago, the underlying basis of those reactions at the molecular level was preferentially selected. Important traits like plant structural traits (plant height, branching, and canopy traits with significant agronomic traits) may be identified using

molecular techniques. For several years, breeders are modifying these architectural plant features to optimize yield and yield [8].

Many genetic factors regulate these architectural traits of plants (Figure 1).

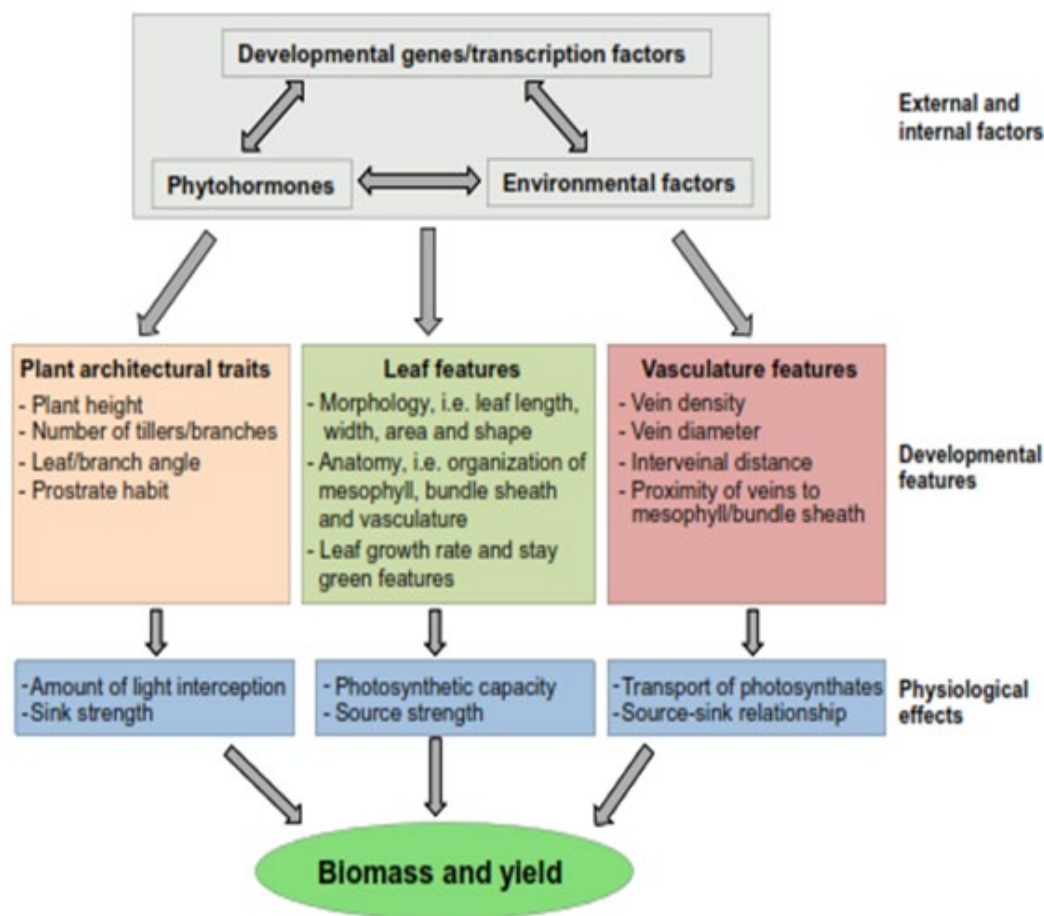


Figure 1: Plant developmental features relevant to crop biomass and yield

1.2. Regulation of Plant Leaf

Leaf characteristics such as shape, size, and thickness are one of the important morphological characteristics that directly affect crop yield [9]. Many recent reviews discuss the mechanisms of the most genetic mechanisms that support leaf events and morphogenesis in both the same coil and bidirectional plants and may increase crop yields. For example, the YABBY PLETHORA (PLT) and JAGGED (JAG) genes actively regulate leaf and leaf plate enlargement within Arabidopsis [10]. The YAB3 gene is involved in stimulating rice leaf growth [11]. In addition, the WUSCHEL-RELATED HOMEBOX (WOX) gene is a gene used to influence leaf size and promote photosynthesis. The rice NARROWLEAF genes (NAL1, NAL2, and NAL3) belonging to the WOX family are associated with increased leaf width through regulation of the cell cycle in early leaf primitive development [12,13].

Many of the above transcription factors affect leaf shape and size by regulating the cell cycle or cell proliferation, making these cellular processes attractive targets manipulated to regulate leaf prop-

erties. The “cell cycle arrest front” is the boundary within the leaf-let primitive where the transition from proliferation to expansion occurs and is important for leaf size regulation. However, further research is needed on its importance in crops and its potential use as a developmental marker of photosynthesis in crops.

1.3. Unraveling miRNA Regulation

MicroRNAs (miRNAs) can be small, unencoded RNAs of nearly 21 nucleotides that regulate gene expression at post-transcriptional levels [14,15]. microRNAs and retargeting genes are widely used to improve the agricultural traits of crops by the expressive master of microRNA genes [16,17]. The relationship of miRNAs between miRNA targets allows users to obtain expressive genes involved in similar biological processes and functions of plants [18,19]. In their study, scrutinize gene regulatory networks and identify key regulatory microRNAs in plant development, they have developed an entirely new algorithm called Differential Edge Transformation (DT) (see Figures 1, 2, and 3).

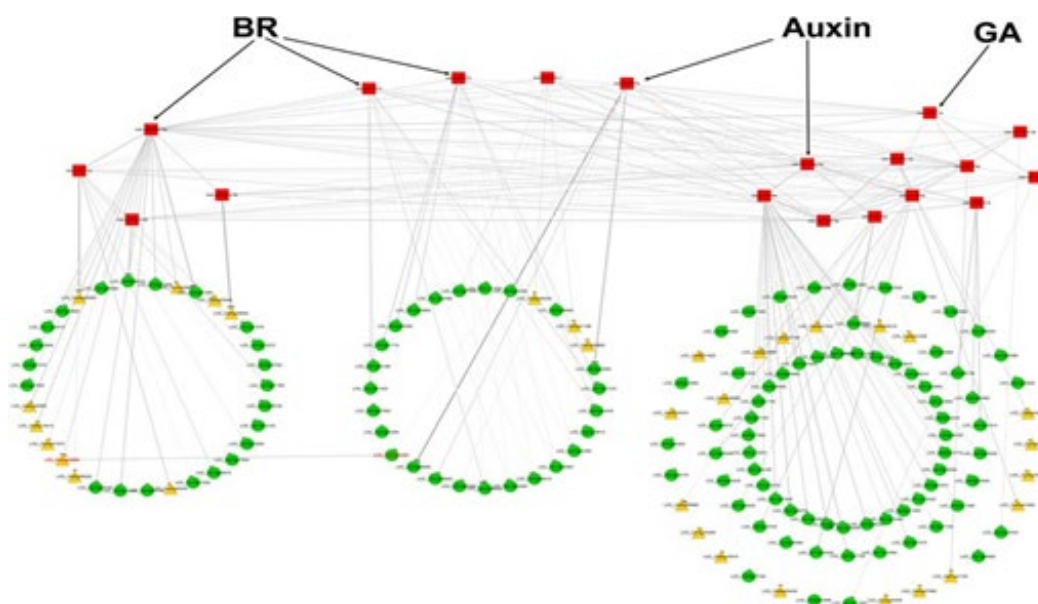


Figure 2; Co-expression of miRNAs identified by differential edge-like transformation target genes for high yields during tillering of rice. Reported yields of miRNAs and their confirmed targets BR (brassinosteroids), auxins and GAs, gibberellins.

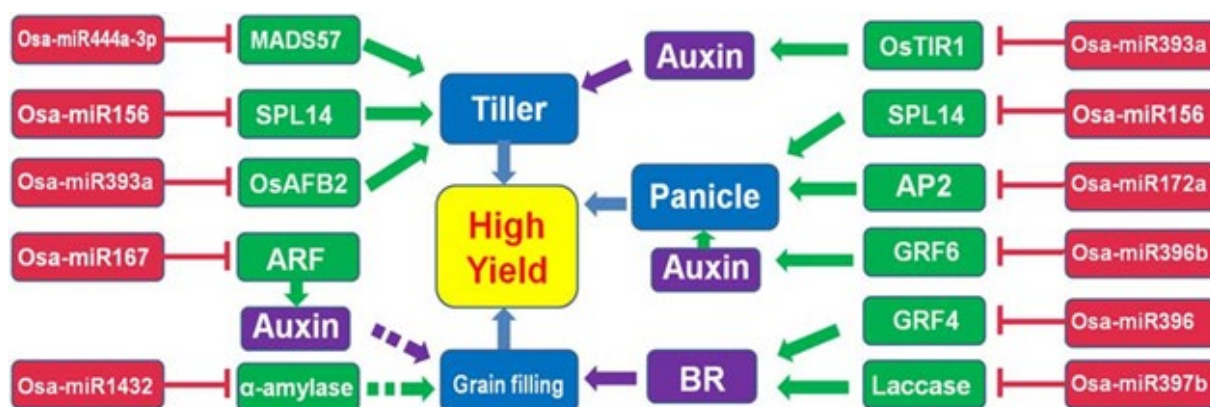


Figure 3: The potential regulatory network model of miRNAs for yields at three stages: tillering, panicle branching, and grain filling in rice. Solid and dashed arrows are the verified and predicted regulatory relationships, respectively.

Ongoing studies show that miRNAs are involved in the regulation of many genes involved in plant structure. For example, Os-miR397 regulates overall yield by controlling grain number and branching per ear through downregulation of its target gene, Os-LAC [19]. miRNAs also regulate other SPL genes that regulate panicle morphology, grain size, and plant structure [20,21]. The molecular properties of the gene indicate that multiple genes con-

tribute to crop yield. For example, the OsNAC2 gene appears to play an important role in the development of rice yields. Typically, revealed that plants overexpressing OsNAC2 under the control of the CaMV35S promoter showed an increase in rice yield [22]. Extensive study of the role of genes in fortifying crops for better yields (see Table 1).

Findings/knowledge	Author(s)/year
Regulation of gene expression in higher plants	Kuhlemeier et al. /1987
Genetic strategies for improving crop yields	Bailey-Serres et al. /2019
Transcriptome analysis and crop improvement (A review)	Dunwell et al. /2001
Crop improvement in the 21st century	Mifflin/(2000
Wild barley: a source of genes for crop improvement in the 21st century?	Ellis et al./2000
Molecular plant breeding as the basis for crop improvement in the 21st century. Plant physiology	Moose & Mumm /2008
Refocus on crop improvement for the changing climatic conditions of the 21st century.	Mba et al./2012
Application of biotechnology to improve crop: prospects and limits	Sharma et al. /2002
Farmers' participatory varietal selection: A sustainable crop improvement approach for the 21st century	Singh et al. /2014
Mutation breeding for crop improvement	Kharkwal et al. /2004
Strategies for improving crops against salt and drought stress: overview	Athar & Ashraf/2009
Shifts in African crop climates by 2050, and the implications for crop improvement and genetic resources conservation	Burke et al./2009
Tissue culture-derived variation in crop improvement. Euphytica	Jain /2001
Improved yields for future agricultural systems. Improving harvest for sustainable agriculture	Francis & Callaway/ 1993
30 underutilized crops: trends, challenges, and opportunities in the 21st century. Managing plant genetic diversity	Padulosi et al/2002
Population genetics of genome-based methods for crop improvement	Hamblin et al. /2011

Table 1: Knowledge, mechanisms mechanism, and role of gene regulation on crop improvement improvements

The molecular properties of the genes are too necessary to analyze the genes of the plant structure used to achieve better yields [23].

2. Conclusion

Crop yield is highly dependent on plant growth and growth. Some developmental traits of the plant, such as general plant structure, leaf traits, and vascular structure, are the most important traits that determine the total yield of the crop. Therefore, enhancing such developmental characteristics has great potential for increasing yields. Regulation of plant genes for plant developmental traits and advances in genomics are revealing the ability to decipher the genes and gene regulatory networks that underlie the developmental traits that are important to agriculture. Combined with an effective approach in the field of phenomenology, this effectively controls the signs of plant growth and increases yields [24-40]. We hope that these approaches will greatly improve our understanding of the genetic basis of domestication of plants and identify additional factors that are targeted for crop improvement. Finally, it should be noted that almost all of these developmental and photosynthetic traits are regulated by many genes and gene regulatory networks. Therefore, modeling, simulation, and systems biology approach also help plant biologists integrate and link genetic networks with developmental and physiological characteristics. This will give you a better understanding of plant growth and improve yields.

Declarations

Ethical Approval and Consent to Participate

Not applicable

Consent for Publication

I as corresponding author declare that we abide by the guideline of the journal and obey publication ethics of the journal.

Availability of Data and Materials

All data were acknowledged and included in this manuscript.

Funding

There is no Funding

Competing Interests

I, the corresponding author verified that there is no competing interest in this manuscript.

Authors' Contributions

Both authors contributed equally

Acknowledgements

N/A

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