

Molecular Mechanism Analysis of Improving the Development of Rice Growth Organs Using Gene Editing Technology

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Abstract Rice (*Oryza sativa*) is one of the most important food crops in the world, and the development of its reproductive organs is crucial for yield and food safety. With the rapid development of gene editing technology, researchers have begun to explore the use of gene editing to improve the development of rice reproductive organs, in order to improve yield and quality. This review aims to explore how gene editing technology can reveal the molecular mechanisms of rice reproductive organ development and explore its potential applications to promote sustainable agriculture and food safety. This study explores the importance of rice as a pillar crop of global food security, as well as the key impact of reproductive organ development on its yield and quality. Then, we introduced the basic principles of gene editing technology, including the working principle of CRISPR-Cas9 technology and the design of RNA guided sequences. Next, we delved into the molecular mechanisms of rice reproductive organs, including the development of roots, leaves, and panicles, and their importance in nutrient absorption, photosynthesis, and seed yield.

Keywords Rice (*Oryza sativa*); Reproductive organ development; Gene editing technology; Sustainable agriculture; Food safety

Rice (*Oryza sativa*), as a primary crop, plays a crucial role in global food security and sustainable agricultural development. The development of its reproductive organs significantly impacts both yield and food quality, making it a critical area in need of exploration within the scientific community. In recent years, with the emergence of gene editing technology, researchers have actively begun to investigate the use of this technology to enhance the development of rice reproductive organs. This is done with the expectation of increasing its yield and quality, ultimately advancing global food security and sustainable agriculture (Zhao et al., 2022).

Globally, rice holds a prominent position as a staple crop, serving as a primary source of food in numerous countries and regions. It directly affects the livelihoods and diets of billions of people. Therefore, understanding the development of rice reproductive organs is of paramount importance in addressing global food security issues. Based on existing research, the development of rice roots, leaves, and panicles has a direct impact on its nutrient uptake, photosynthetic efficiency, as well as the ultimate yield and quality of seeds. This intricate relationship underscores the critical need for in-depth investigation into rice reproductive organ development, providing essential insights for optimizing cultivation conditions and enhancing crop yield.

This study is driven by the pursuit of understanding the molecular mechanisms underlying rice reproductive organ development. Gene editing technology, particularly CRISPR-Cas9, is regarded as a groundbreaking tool that offers a potential avenue for genome improvement in rice. The primary objective of this research is to explore how gene editing technology can unveil the molecular mechanisms of rice reproductive organ development. Our aim is not only to comprehend the developmental processes of rice reproductive organs but also to investigate the potential applications of gene editing technology in optimizing these processes.

Through this research, we aspire to provide a more forward-looking direction for future agricultural development. Rice, as one of the world's critical crops, represents not only an enhancement in individual yield but also a significant step towards combating hunger, promoting agricultural sustainability, and safeguarding global food security. This study aims to offer a comprehensive perspective on the molecular mechanisms of rice reproductive

organ development and the application of gene editing technology in this field. It seeks to chart a course for future research directions and is dedicated to propelling global agriculture towards a more sustainable and innovative trajectory.

1 Importance of Rice Reproductive Organ Development

1.1 Rice as a global staple food crop

Rice is hailed as one of the "world's most crucial staple crops", playing a pivotal role in global food security. It constitutes a significant portion of global food production, meeting the primary dietary needs of billions of people. This study will delve into the significance of rice in global food production and food security, as well as how its status as a major crop propels research and improvement in the development of rice reproductive organs.

Rice, as one of the world's primary staple crops, particularly in Asian regions, serves as a primary food source for many countries, providing a major source of carbohydrates and energy for billions of people. Its seeds, known as rice grains, are an indispensable part of human diet, offering a primary source of energy and nutrients.

Rice not only provides the energy required by humans but also offers a variety of nutrients such as proteins, vitamins, and minerals. It serves as the primary food source for millions of households. Therefore, research on rice yield and quality is of paramount importance for global food security (Zhao et al., 2020). Both yield and quality have been subjects of continuous scrutiny, as agricultural scientists have been actively seeking methods to enhance rice yield and quality. The development of rice reproductive organs is one of the key factors in achieving this goal, as it directly influences rice yield and quality.

This study will delve deeply into the development of rice reproductive organs, including roots, leaves, and panicles, to unveil their molecular mechanisms and explore how gene editing technology can be harnessed to enhance these processes. The goal is to boost rice yield and quality. These research endeavors will contribute to addressing global food security challenges and ensuring that rice continues to play its pivotal role in the global food supply chain.

1.2 Impact of reproductive organ development on yield and quality

Rice reproductive organs encompass roots, leaves, and panicles, serving as the foundation for rice growth and reproduction. The developmental status of these reproductive organs directly influences both rice yield and quality (Figure 1). Investigating the impact of rice reproductive organ development on yield and quality is crucial for gaining a deeper understanding of why researching and improving these processes is of paramount importance.

The root system of rice is the primary organ for absorbing moisture and nutrients from the soil, and the developmental status of the root system directly affects the efficiency of rice in absorbing water and nutrients. A healthy and well-developed root system can more effectively absorb nutrients such as nitrogen, phosphorus, and potassium, thereby enhancing rice growth and yield. Therefore, researching and improving the development of the root system is of paramount significance in increasing rice yield.

Leaves, as the primary organs for photosynthesis in plants, convert light energy into chemical energy, supporting plant growth and development. The number, size, and health of leaves affect the efficiency of photosynthesis, consequently impacting rice yield (Adachi et al., 2019). Normal leaf development and the stability of photosynthesis are equally crucial in increasing rice yield and improving quality (Figure 1), and these factors should not be overlooked.

Panicles serve as the reproductive organs of rice, responsible for seed production and dissemination. The developmental status of panicles directly determines rice seed yield. Healthy panicles and effective pollen dissemination are crucial factors for high-yielding rice (Figure 1). Researching and improving the development of panicles significantly impact the ultimate yield and quality of rice.

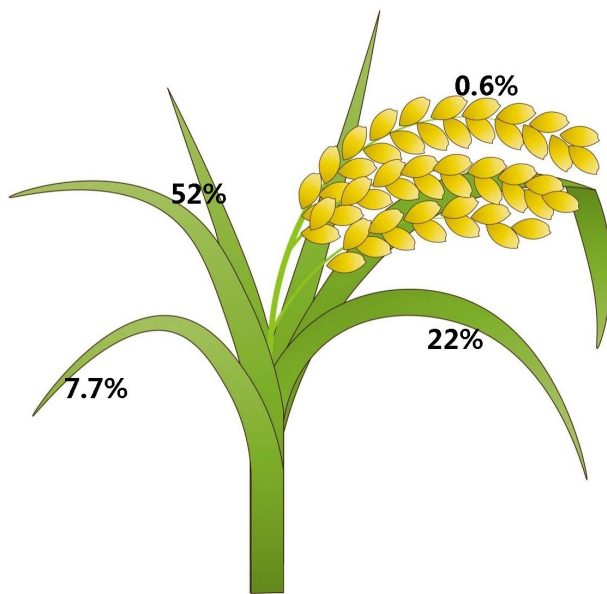


Figure 1 Contribution rate of different rice leaves to yield

The reproductive organs of rice, including roots, leaves, and panicles, play a crucial role in both yield and quality. Understanding the developmental processes of these reproductive organs and their impact on rice growth is pivotal for achieving sustainable agriculture and food security. Through in-depth research into the molecular mechanisms of these organs and the application of gene editing technology, there is hope for further enhancing rice yield and quality to meet global food demands.

1.3 The potential of gene editing technology in rice improvement

Gene editing technology has become a powerful tool for improving crops, including rice. The potential of gene editing technology in rice improvement includes the application of CRISPR-Cas9 technology and its contribution to improving rice yield and quality. CRISPR-Cas9 technology is a revolutionary gene editing tool that allows scientists to accurately modify rice genes. This technology achieves gene editing by designing specific RNA guidance sequences to guide Cas9 proteins onto target genes (Li et al., 2020). In rice, CRISPR-Cas9 technology has been widely used to modify genes related to reproductive organ development. Scientists have successfully edited genes related to root development to improve the root structure and nutrient absorption efficiency of rice. In addition, the development of leaves and ears has also been studied and improved by CRISPR-Cas9 technology.

Gene editing technology provides a new avenue for improving rice yield and quality. By precisely editing genes associated with reproductive organ development, we can enhance the root's nutrient absorption capacity, improve photosynthetic efficiency, increase panicle yield, and adjust rice's resilience to cope with climate change and pests. These improvements will contribute to increasing rice yield, providing more high-quality food to meet global food demands.

While gene editing technology holds tremendous potential for rice improvement, it also faces certain technical challenges and ethical concerns. These include issues related to editing efficiency, uncertainty in the effects of edits, ecological risks, as well as food safety and ethical considerations. Scientists and policymakers need to collaborate to address these challenges to ensure the safe and sustainable application of gene editing technology.

Throughout history, gene editing technology has provided unprecedented opportunities for rice improvement by precisely editing genes related to reproductive organ development to enhance yield and quality. However, the application of this technology requires careful consideration of its potential challenges and risks to ensure sustainable agriculture and food security. Future research and practices will continue to uncover the potential of gene editing technology in rice improvement, offering new pathways to address global food challenges.

2 Gene Editing Technology Fundamentals

2.1 Introduction of CRISPR-Cas9 technology

Clustered Regularly Inter-spaced Short Palindromic Repeats (CRISPR) is a natural immune system found in bacteria and archaea. CRISPR-Cas9 technology is a revolutionary gene editing tool that has achieved significant success in plant improvement. This system can record and identify foreign DNA and cut and repair it using Cas proteins. Scientists have discovered that, by utilizing the CRISPR system, they can precisely cut specific DNA sequences by designing specific RNA guide sequences to guide the Cas9 protein. This technology allows scientists to precisely edit target genes, enabling the addition, deletion, or replacement of specific gene segments (Romero and Gatica-Arias, 2019).

In rice improvement, CRISPR-Cas9 technology has been widely applied. Scientists can design appropriate RNA guide sequences to direct the Cas9 protein into rice genes, enabling the editing of target genes. This editing can encompass genes related to the development of reproductive organs such as roots, leaves, and panicles. Some studies suggest that by editing genes related to root development, improvements in the root structure and nutrient absorption efficiency of rice can be achieved. Simultaneously, editing genes in leaves and panicles also holds the potential to enhance photosynthetic efficiency and seed yield in rice.

The high precision and customization of CRISPR-Cas9 technology make it a crucial tool in rice improvement. It provides scientists with a pathway to enhance rice yield, quality, and resilience to adapt to different growth environments and climate conditions. Although the technology still faces some challenges and ethical concerns, it undoubtedly offers unprecedented opportunities for rice improvement, with the potential to drive global food security and sustainable agriculture.

2.2 Design and selection of RNA guide sequences

RNA guide sequences play a pivotal role in CRISPR-Cas9 technology, serving as guides to direct the Cas9 protein to specific DNA sequences within the target gene. Designing and selecting appropriate RNA guide sequences are of utmost importance for successful gene editing. The design of RNA guide sequences should adhere to the following principles to ensure precise gene editing: RNA guide sequences must exhibit a high degree of matching with the specific DNA sequence of the target gene to prevent unintended editing of non-target genes.

At the same time, it is essential to avoid any matching of the RNA guide sequences with non-target DNA sequences in the rice genome to reduce unnecessary gene editing. RNA guide sequences should not align with repetitive sequences within the rice genome to prevent misguided editing. Selecting appropriate RNA guide sequences typically involves a comprehensive analysis of the target gene, and specialized computational tools can aid in the screening of the best RNA guide sequences (Usman et al., 2020).

RNA guide sequences should be selected in close proximity to the target gene to enhance editing efficiency. The specificity of RNA guide sequences is of paramount importance, ensuring that they do not match to other positions within the rice genome. An optimal GC content contributes to the stability and efficiency of RNA guide sequences. Computational tools can assist in predicting the editing efficiency of different RNA guide sequences, allowing scientists to choose the most efficient sequences.

Once suitable RNA guide sequences are designed and selected, they are used in conjunction with the Cas9 protein within the CRISPR-Cas9 system to achieve precise editing of rice genes. Through thoughtful RNA guide sequence design and selection, scientific research can further enhance the success rate of CRISPR-Cas9 technology in rice improvement, with the potential to contribute to global food security and sustainable agriculture.

2.3 Mechanism and applications of gene editing

Gene editing technology is a method for precise genetic improvement by modifying an organism's DNA sequence. In the study of rice reproductive organ development, gene editing technology has become a crucial tool, aiding in the in-depth understanding of developmental mechanisms of reproductive organs and enhancing the traits of rice.

The core mechanism of gene editing technology typically involves the following key components. The Cas protein within the CRISPR-Cas9 system is an RNA-guided endonuclease responsible for cutting the target DNA molecule. In rice research, Cas9 is the most commonly used Cas protein. The RNA guide sequence is a specific RNA sequence that pairs with the complementary DNA sequence of the target gene. It can guide the Cas protein to a specific gene locus, thereby enabling gene editing.

Gene editing typically involves DNA repair mechanisms, including non-homologous end joining (NHEJ) and homology-directed repair (HDR). NHEJ often leads to small insertions or deletions, while HDR can achieve more precise gene editing. By guiding Cas9 and the RNA sequence to the target gene, gene knockout can be achieved. In the study of rice reproductive organ development, this can assist scientists in determining the function of specific genes in reproductive organ development (Chang et al., 2009).

Gene editing technology can also be used to repair damaged genes. Scientists can provide repair templates to replace the damaged DNA sequence through the HDR mechanism, thereby restoring the gene. In rice research, scientists can also utilize gene editing to introduce foreign genes, enhancing rice's resilience to stress or improving traits. This can be achieved through the HDR mechanism.

While gene editing technology holds vast potential for research on rice reproductive organ development, it still faces some challenges. However, as the technology continues to advance, and with the increasing global recognition of gene editing, we can expect more innovative applications of this technology in the study of rice reproductive organ development, aimed at improving rice yield, quality, and resilience.

3 Molecular Mechanism Analysis of the Development of Rice Growth Organs

3.1 Molecular mechanisms of rice root development and nutrient uptake

The roots of rice are a critical component of the reproductive organs, providing structural support and stability while also being responsible for nutrient and water absorption, which is vital for rice growth and development. The root tip comprises a meristematic tissue that continuously generates new root cells, and the activity in this region is regulated by growth hormones such as gibberellins and auxins. The activity of the meristematic tissue sustains root growth, allowing it to continuously extend downward. The outer surface of the roots is covered with root hairs, which are crucial sites for nutrient absorption. The development of root hairs is a highly precise process, involving reshaping and growth through cell wall modification. Gene families such as Root Hair Defective 6 (RHD6) and RHD6-LIKE play pivotal roles in root hair development (Yang et al., 2023).

The process of nutrient uptake by the roots is complex and involves several key nutrients, including nitrogen, phosphorus, and potassium. There are nitrogen uptake organs in the roots, such as amino acid transport proteins, which assist rice in absorbing nitrogen sources from the soil. Nitrogen is a critical component for plant growth and protein synthesis, making it essential for the healthy growth of rice. Phosphorus is another vital nutrient for rice, and it is absorbed through specialized cell membrane channels in the roots. Improving root development can enhance the efficiency of phosphorus uptake, thereby improving rice yield and quality. Potassium is an essential element for plant growth and development, and it is absorbed through potassium channels on the cell membrane of the roots. Potassium helps maintain cell osmotic pressure and enzyme activity.

Understanding the molecular mechanisms of root organ development and nutrient absorption contributes to scientists' efforts to improve nutrient uptake efficiency in rice, thereby increasing yield and quality. Through gene editing technology, root development and nutrient absorption processes can be precisely regulated, making a valuable contribution to global food security and sustainable agriculture.

3.2 Leaf development and photosynthesis

Leaves are the primary organs responsible for photosynthesis in rice plants, playing a crucial role in rice growth and development. Leaf development is a complex process influenced by multiple genes and regulatory factors. It begins with the formation of leaf primordia, during which a series of gene families, such as MADS-box and TCP,

play pivotal roles in regulating early differentiation and development of leaves. The synthesis of chlorophyll in leaves is an integral part of leaf development. Chlorophyll is a key molecule in photosynthesis, enabling rice plants to convert light energy into chemical energy (Wang et al., 2020).

The surface of leaves is covered with chlorophyll, and the interior of leaves contains chloroplasts that house enzymes and other molecules required for photosynthesis. These physiological structures make leaves the focal point for photosynthesis. Photosynthesis comprises two primary stages: the light reactions and the carbon reactions. In the light reactions, leaves absorb light energy, convert it into electron energy, and produce oxygen (Figure 2). In the carbon reactions, leaves utilize light energy to convert carbon dioxide into carbohydrates such as glucose. The process of photosynthesis is regulated by factors such as light intensity, temperature, and moisture. Plants can adjust the influx and efflux of carbon dioxide by opening and closing stomata to adapt to varying environmental conditions.

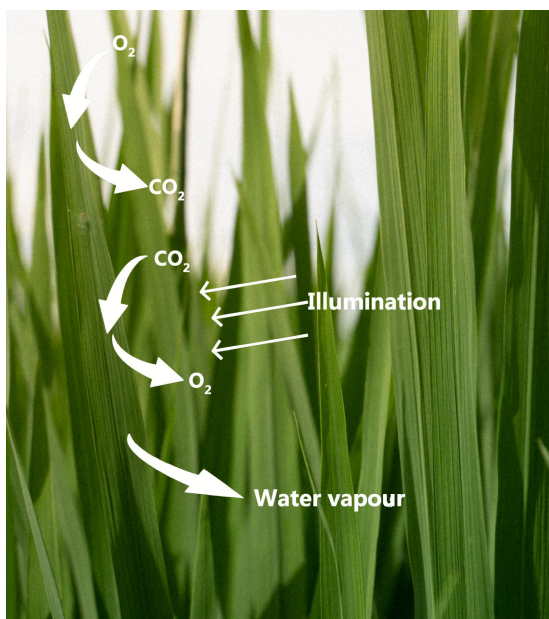


Figure 2 Photosynthesis process of rice leaves

The development of leaves and photosynthesis are closely intertwined, and together, they impact the growth and yield of rice. Through in-depth research into the developmental processes of leaves and the molecular mechanisms of photosynthesis, scientists can seek methods to enhance rice photosynthetic efficiency, increase resilience to environmental stress, and improve yield.

The development of the panicle, a crucial organ for seed production in rice, is closely related to the ultimate seed yield. Understanding the developmental mechanisms of the panicle is of significant importance for improving rice yield and quality. Panicle development begins with the differentiation of axillary buds, which will gradually form primary branches and tillers, generating multiple spikelets. Each spikelet comprises a series of florets that arrange themselves into a sequence at the apex of the panicle. Spikelet formation involves cell division and differentiation, regulated by growth hormones. The florets within each spikelet develop consecutively, including the formation of male and female reproductive organs and embryo sac development.

The number of florets within the panicle determines the final seed quantity, and increasing seed yield can be achieved by promoting the differentiation of axillary buds and floret development. In addition to quantity, the quality of florets is also of paramount importance. Factors such as the size, shape, and quality of florets within the panicle can influence the ultimate seed yield and quality. The development of reproductive organs in florets, the production of pollen, as well as the pollination process also have an impact on seed yield.

Understanding the developmental mechanisms of panicles and their relationship with seed yield is crucial for improving rice production. Through in-depth research into these molecular mechanisms, scientists can take measures to increase the number of florets and improve floret quality, thereby enhancing rice seed yield. This has the potential to meet global food demand and is of significant importance for sustainable agriculture and food security.

4 Application of Gene Editing Technology in Improving Rice Organ Development

4.1 Gene editing cases in root development

The application of gene editing technology in rice root development has yielded some promising cases, providing new possibilities for increasing rice yield and adapting to different soil conditions.

A study conducted by the Chinese Academy of Agricultural Sciences, scientists successfully edited genes related to root growth and branching in rice. By editing specific genes, they achieved growth promotion of rice roots and increased branching. These improved rice varieties have shown stronger drought resistance and better nutrient absorption ability, thereby increasing yield.

Another gene editing case involves increasing the ability of rice roots to absorb specific nutrients such as phosphorus. Scientists used CRISPR-Cas9 technology to edit phosphorus absorption related genes in rice, making it more effective in absorbing phosphorus from soil. This improvement helps to reduce dependence on fertilizers, reduce environmental pollution, and increase rice yield (Li et al., 2020).

The root system is not only used to absorb nutrients and water, but also to combat parasitic pests in the roots. Gene editing technology has the potential in the development and functional regulation of rice roots, and precise editing of specific genes can improve rice growth and adaptability. Scientists use gene editing technology to enhance the resistance of rice roots to pests. By editing genes related to plant defense mechanisms, rice varieties with stronger insect resistance have been created, reducing the use of pesticides and improving yield and quality.

4.2 A case study of gene editing in leaf development

Gene editing technology has also produced some real cases in the field of rice leaf development, which provide new ways to increase rice yield and improve leaf characteristics. In a study, scientists used CRISPR-Cas9 technology to edit genes related to chlorophyll synthesis in rice. They successfully increased the content of chlorophyll in leaves, enabling them to better absorb light energy, thereby improving the photosynthetic efficiency of rice. This improvement helps to increase the growth rate and yield of rice.

The size and shape of leaves have a significant impact on physiological processes such as photosynthesis and water evaporation. Scientists have used gene editing techniques to regulate genes related to cell division and expansion in rice leaves, altering their morphology. It is precisely because of this change that larger or smaller leaves have been successfully created to adapt to different growth environments (Zhang et al., 2020).

In addition to influencing the intrinsic characteristics of leaves, gene editing can also be employed to enhance the resistance of rice leaves to diseases and pests. By editing genes related to plant defense mechanisms, rice leaves can better withstand pathogen attacks, reducing damage from diseases and pests and further improving yield and quality. Therefore, gene editing technology holds the potential for applications in rice leaf development and trait regulation, allowing for the precise editing of genes associated with leaves to enhance photosynthetic efficiency, disease resistance, and morphology, ultimately increasing rice yield and adaptability.

4.3 Gene editing cases of ear development

Gene editing technology has also yielded some real cases in the field of rice panicle development. Scientists have used CRISPR-Cas9 technology to edit genes related to rice panicle differentiation and tillering. They successfully increased the efficiency of tiller bud differentiation, promoting the formation of more tillers in rice plants, allowing them to produce more panicles, potentially increasing the number of seeds and thus improving yield.

Another case involves extending the growth period of panicle development. By editing genes in rice related to the growth cycle of panicles, scientists have enabled them to develop and produce more panicles over an extended period. Through comparative observations, it has been found that rice produces more florets, directly increasing yield and the number of seeds (Wang et al., 2020).

Panicle development is influenced by environmental factors such as drought or pest infestations, and gene editing technology has enhanced rice plants' resistance to stressors. They have edited genes related to plant stress resistance, allowing panicles to better withstand adverse conditions, protecting floret development, and thereby increasing yield. The potential of gene editing technology in regulating panicle development and function in rice, through the precise editing of genes related to panicle development, can improve rice growth and adaptability.

5 Future Outlook

Rice is one of the most crucial global staple crops, essential for human food security. However, with the world's increasing population and the impact of climate change, the demand for rice yield and quality continues to rise, compelling humans to seek new ways to improve rice reproductive organ development. Gene editing technology has shown tremendous potential, providing new opportunities for genetic improvement in rice.

With the continuous advancement of gene editing technology, researchers can precisely edit genes related to rice reproductive organ development. This will contribute to more targeted genetic improvements to enhance the characteristics of rice roots, leaves, and panicles. By precisely editing target genes, humans can achieve higher yields, better resilience, and improved quality.

Climate change has brought about challenges such as drought, saline-alkali soils, and pests. Gene editing technology can be used to develop rice varieties with enhanced resilience. By editing genes related to stress responses, more stress-tolerant rice can be created to adapt to various environmental conditions, ensuring stable food production.

Gene editing technology and traditional breeding methods can complement each other, offering more options for rice genetic improvement. Traditional breeding often takes a long time to develop new varieties, whereas gene editing can expedite this process. In the future, we can utilize gene editing to accelerate the breeding process while retaining other beneficial traits of rice (Zafar et al., 2020).

With the continuous growth of the global population, meeting the demand for food has become a crucial issue. Gene editing technology offers opportunities to increase rice yield, improve quality, and adapt to different soil conditions. This contributes to ensuring food supply, maintaining food security, reducing reliance on fertilizers and pesticides, decreasing agriculture's adverse environmental impact, and promoting the development of sustainable agriculture.

As the application of gene editing technology progresses, legal, ethical, and safety issues will also garner widespread attention. Ensuring the safety and sustainability of gene editing is a critical challenge. The international community needs to establish relevant regulations and ethical guidelines to guide the application of gene editing technology, while also ensuring food safety and environmental sustainability.

Gene editing technology offers limitless possibilities for genetic improvement in rice organ development. In the future, we can expect more innovations and real-world cases to ensure increased rice yield, improved quality, and the maintenance of food security. This requires cooperation from the international community and the relentless efforts of scientists to achieve better genetic improvements in rice and sustainable agriculture.

References

- Adachi S., Yamamoto T., Nakae T., Yamashita M., Uchida M., Karimata R., Ichihara N., Soda K., Ochiai T., Risako Ao, Otsuka C., Nakano R., Takai T., Ikka T., Kondo K., Ueda T., Ookawa T., and Hirasawa T., 2019, Genetic architecture of leaf photosynthesis in rice revealed by different types of reciprocal map populations, *Journal of Experimental Botany*, 70(19): 5131-5144.
<https://doi.org/10.1093/jxb/erz303>

- Chang Y., Gong L., Yuan W., Li X.W., Chen G.X., Li X.H., Zhang Q.F., and Wu C.Y., 2009, Replication protein A (RPA1a) is required for meiotic and somatic DNA repair but is dispensable for DNA replication and homologous recombination in rice, *Plant physiology*, 151(4): 2162-2173.
<https://doi.org/10.1104/pp.109.142877>
- Ding S.L., Liu C.L., and Qian Q., 2019, Research progress on genetic of rice root, *Zhongguo Daomi (China Rice)*, 25(5): 24.
- Li J., Yokosho K., Liu S., Cao H.R., Yamaji N., Zhu X.G., Liao H., Ma J.F., and Chen Z.C., 2020, Diel magnesium fluctuations in chloroplasts contribute to photosynthesis in rice, *Nature Plants*, 6(7): 848-859.
<https://doi.org/10.1038/s41477-020-0686-3>
- Li S.L., Zhen H.Y., and Wang L., 2020, Application and prospect of gene editing technology in crop breeding, *Shengwu Jishu Tongbao (Biotechnology Bulletin)*, 36(11): 209.
- Romero F.M., and Gatica-Arias A., 2019, CRISPR/Cas9: development and application in rice breeding, *Rice Science*, 26(5): 265-281.
<https://doi.org/10.1016/j.rsci.2019.08.001>
- Usman B., Nawaz G., Zhao N., Liao S.Y., Liu Y.G., and Li R.B., 2020, Precise editing of the OsPYL9 gene by RNA-guided Cas9 nuclease confers enhanced drought tolerance and grain yield in rice (*Oryza sativa* L.) by regulating circadian rhythm and abiotic stress responsive proteins, *International Journal of Molecular Sciences*, 21(21): 7854.
<https://doi.org/10.3390/ijms21217854>
- Wang C., Chen S., Dong Y., Ren R.J., Chen D.F., and Chen X.W., 2020, Chloroplastic *Os3BGLu6* contributes significantly to cellular ABA pools and impacts drought tolerance and photosynthesis in rice, *New Phytologist*, 226(4): 1042-1054.
<https://doi.org/10.1111/nph.16416>
- Wang F., Han T., Song Q., Ye W.X., Song X.G., Chu J.F., Li J.Y., and Chen J., 2020, The rice circadian clock regulates tiller growth and panicle development through strigolactone signaling and sugar sensing, *Plant Cell*, 32(10): 3124-3138.
<https://doi.org/10.1105/tpc.20.00289>
- Wang T., Li Y., Song S., Qiu M.D., Zhang L.C., Li C.X., Dong H., Li L., Wang J.L., and Li L., 2021, EMBRYO SAC DEVELOPMENT 1 affects seed setting rate in rice by controlling embryo sac development, *Plant Physiology*, 186(2): 1060-1073.
<https://doi.org/10.1093/plphys/kiab106>
- Yang S., Xu N., Chen N., Qi J.X., Abdul S., Wu J.Y., Liu Y.H., Huang L.L., Liu B.H., and Gan Y.B., 2023, OsUGE1 is directly targeted by OsGRF6 to regulate root hair length in rice, *Theoretical and Applied Genetics*, 136(5): 108.
<https://doi.org/10.1007/s00122-023-04356-4>
- Zafar K., Sedeek K.E.M., Rao G.S., Shahid M.M., and Mahfouz M., 2020, Genome editing technologies for rice improvement: progress, prospects, and safety concerns, *Frontiers in Genome Editing*, 2: 5.
<https://doi.org/10.3389/fgeed.2020.00005>
- Zhang Y., Zhang Y., and Xie K.B., 2020, Evaluation of CRISPR/Cas12a-based DNA detection for fast pathogen diagnosis and GMO test in rice, *Molecular Breeding*, 40: 1-12.
<https://doi.org/10.1007/s11032-019-1092-2>
- Zhao M., Lin Y., Chen H., 2020, Improving nutritional quality of rice for human health, *Theoretical and Applied Genetics*, 133: 1397-1413.
<https://doi.org/10.1007/s00122-019-03530-x>
- Zhao Z., Wang C., Yu X., and Wang J.M., 2022, Auxin regulates source-sink carbohydrate partitioning and reproductive organ development in rice, *Proceedings of the National Academy of Sciences*, 119(36): e2121671119.
<https://doi.org/10.1073/pnas.2121671119>