

APPLICATIONS OF THE CRISPR/CAS9 TECHNIQUE IN MAIZE AND WHEAT BREEDING

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Abstract

Maize and wheat are some of the world's most important food crops, so their breeding programs are important for global food security. Genome editing techniques (the latest advancement in genetics) are not replacements for conventional breeding techniques, they are new methods and innovative, which promote agricultural crop breeding programs and offers unprecedented solutions to food insecurity. Among these techniques, CRISPR-Cas9 is considered a more effective tool for genome editing due to its low cost and simplicity. This paper summarizes recent applications of the CRISPR/Cas9 technique in maize and wheat breeding. The implementation of this technique allows the production of non-transgenic crops with high yield under different environmental stresses promoting sustainable agriculture.

Key words: CRISPR/Cas9, genome editing, maize, wheat

INTRODUCTION

The world's population rapidly increased and continues to do so, estimating that in 2050 it will reach 10 billion, therefore a 60-100% increase in world food production will be necessary (Dhankher and Foyer, 2018), while the agricultural area on per capita is decreasing every year (Ritchie and Roser, 2019). This situation is aggravated by climate changes that affect agricultural systems and food security. Therefore, obtaining new crop varieties (with high yield and resistant to biotic and abiotic stress) is the main objective of plant breeding (Bonea, 2020). Maize and wheat are some of the most important crops used as food, feed and industrial raw material that ensure food security for billions of people (Ștefan and Constantinescu, 2011; Constantinescu and Olaru, 2017; Bonea, 2020; Constantinescu et al., 2021). Due to their worldwide importance, these crops have become a target for genetic

improvement. According to the Food and Agriculture Organization, in 2020/2021 of the total global production of maize (1483.2 million tonnes), 871.1 million tonnes were utilized as feed, 393.6 million tonnes to other non-food uses and 223 million tonnes as food, and of the total global production of wheat (776.7 million tonnes), 148 million tonnes were utilized as feed, 88.9 million tonnes to other non-food uses and 525.5 million tonnes as food (FAO, 2022).

Farmers and plant breeders have developed varieties of various agricultural plants through conventional technologies, such as selection and propagation of plants with useful traits. Genetic modification of plants (transgenesis) is a modern breeding technique used to improve agricultural crops which consists in modifying the genome of an organism, as a result of introducing new genes (transgenes), or as a result of changing the expression of one/some genes already present in the cell (Bonea, 2013; Urechean

and Bonea, 2017; Bonea, 2021). In 1996, genetically modified crops occupied only 1.7 million hectares, and in 2018 it reached an area of 191.7 million hectares (Bonea, 2022). However, these genetically modified crops have been controversial for various reasons (agricultural policies, insufficient information, public concerns, etc) (Ezezika et al., 2012). Also, concerns regarding the use of certain chemicals (e.g. glyphosate) in combination with herbicide-tolerant genetically modified plants, or the transmission of antibiotic resistance genes, have led to the adoption of very strict regulations on genetically modified plants, making it difficult to produce varieties suitable for current threats (Zaidi et al., 2020).

Recently, new plant breeding technologies have emerged that include genome editing and differ from genetic modification.

According to scientists, gene/genome editing is not "genetic modification" because the method of introducing DNA changes is no different from changes that can occur during conventional reproduction or in nature (Pacher and Puchta, 2017). For example, CRISPR-Cas can be used for precise genetic manipulation without inserting exogenous DNA, such as antibiotic resistance genes, thus eliminating the fear that foreign DNA may be present in the final product (He and Zhao, 2020).

This paper presents a summary of recent CRISPR/Cas9 technique applications in the maize and wheat breeding.

GENOME EDITING

Genome editing includes a series of molecular techniques that allow the induction of directed (targeted) changes in the genomes of organisms.

Four major classes of sequence-specific nucleases (SSNs) are used for genome editing: Meganucleases (MNs); Zinc-Finger Nucleases (ZFNs); TALENs (Transcription Activator-Like Effector Nucleases) and proteins Cas9. These nucleases can be engineered to bind and cleave a specific nucleic acid sequence, introducing double-strand breaks (DSBs) at or near the target site (Pickar-Oliver and Gersbach, 2019).

Figure 1 shows the advantages and disadvantages for each of these genome editing techniques.

CRISPR/CAS9 TECHNIQUE FOR CROP IMPROVEMENT

Clustered Regularly Interspersed Short Palindromic Repeats/CRISPR-associated protein 9 (CRISPR/Cas9) system has two main components: the Cas9 protein which produces DSBs (double-strand break) at a targeted site and a single guide RNA (sgRNA) that identifies a specific DNA sequence thus, when changing the design of sgRNA, numerous desired sites can be targeted (Jinek et al., 2012; Kim and Kim, 2014).

Figure 2 shows the mechanism of genome editing using CRISPR/Cas9 involving DNA unwinding by sgRNA, cutting of gene by Cas9, genome analysis, sgRNA cloning, plants transformation and selection, regeneration, genomic DNA extraction and sequence analysis to confirm the results. Therefore, this process needs no foreign element for editing (Rasheed et al., 2021). The CRISPR/Cas9 technique is a more versatile genome editing technique that has many advantages: low cost, good adaptability, time efficiency, the ability to instantly direct the reproduction of multiple genes, it is also more cost-effective and simpler compared to ZFNs, TALENs and MNs (Hsu et al., 2014; Braatz et al., 2017).

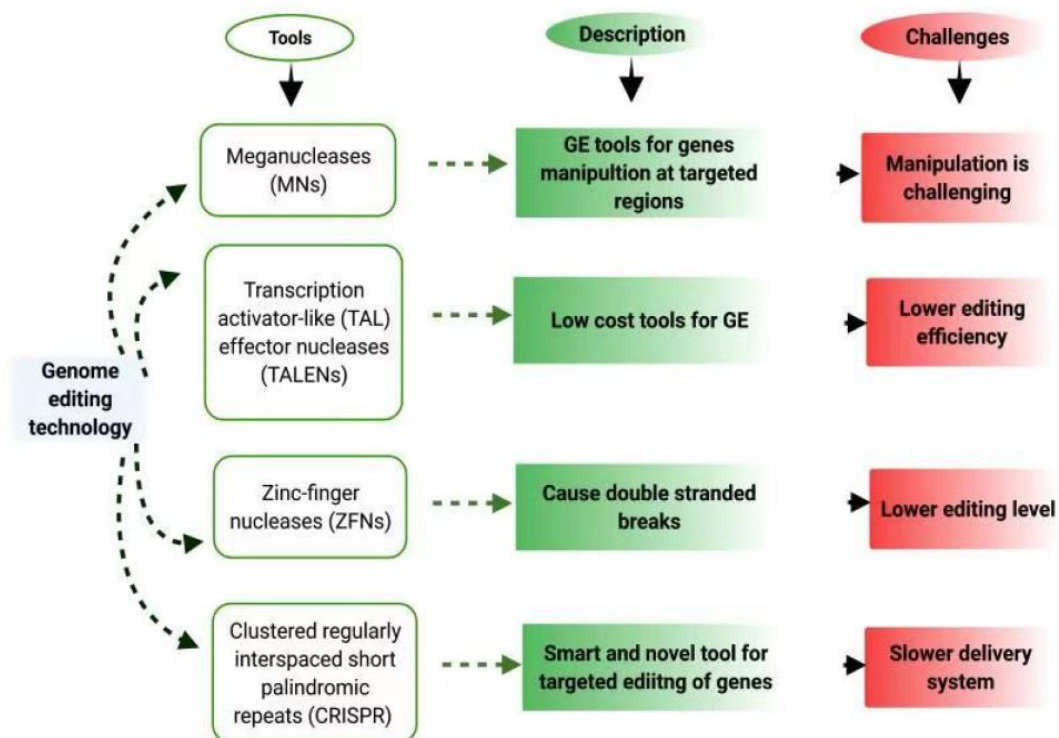


Figure 1. A comparison of genome editing tools regarding their efficiency and limitations

(Source: Rasheed et al., 2021)

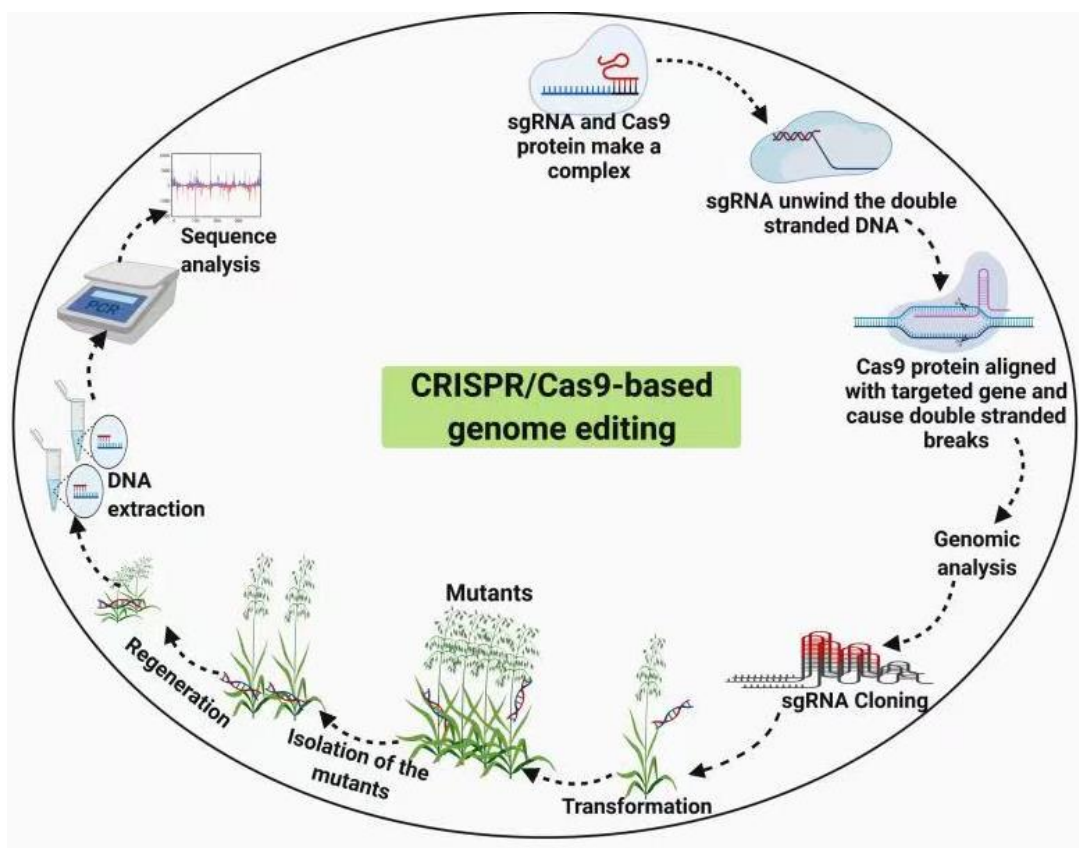


Figure 2. Mechanism of genome editing using CRISPR/Cas9 (Source: Rasheed et al., 2021)

Due to these advantages, CRISPR/Cas9 has been used to improve some traits in many monocot and dicot crops, such as yield, quality, and resistance to biotic and abiotic stresses (Ma and Liu, 2016).

Creating a new approach to manipulating the genomes of living organisms, genome editing has revolutionized the biological sciences.

Many techniques have been developed to improve some traits to crops, but CRISPR/Cas9 is the most useful and latest method (Rasheed et al., 2021).

According to Martin-Laffon et al. (2019), the total number of patents on the CRISPR-Cas9 landscape until 2019 was of 1052, in agriculture was of 374, in industry was of 192 and in medical was of 614.

Zion Market Research firm projected that the CRISPR-Cas9 gene editing market will

reach US\$4.271 billion by 2024, a compound annual growth rate (CAGR) of 36.8%, and Grand View Research firm anticipates the global market for gene editing will reach \$8.1 billion by 2025 (CRS, 2018).

CRISPR/Cas, is a critical gene editing tool that was developed in 2013 by two scientists named Jennifer Doudna and Emmanuelle Charpentier, who demonstrated that CRISPR can be used to modify human genes outside the body (Jinek et al., 2012; Doudna and Charpentier, 2014).

This technique is currently used for various aspects in plant breeding such as yield and quality, disease resistance, drought tolerance.

Table 1 shows some recent applications in maize and wheat breeding.

Table 1. Examples of recent applications of the CRISPR/Cas9 technique to maize and wheat breeding

Species	Trait improved	Technique	Gene(s) edited	References
Maize	grain yield	CRISPR/Cas9	<i>Waxy</i>	Gao et al. (2020)
	grain yield	CRISPR/Cas9	<i>CLE</i>	Liu et al. (2021)
	drought tolerance	CRISPR/Cas9	<i>ARGOS8</i>	Shi et al. (2017)
	herbicide tolerance	CRISPR/Cas9	<i>ZmALS1, ZmALS2</i>	Nuccio et al. (2021)
	salinity tolerance	CRISPR/Cas9	<i>ZmHKT1</i>	Zhang et al. (2018)
	reduced phytic acid	TALENs, CRISPR/Cas9	<i>ZmPDS, ZmIPK1, ZmIPK, ZmMRP4</i>	Liang et al. (2014)
	reduced zein content	CRISPR/Cas9	<i>PRL</i>	Qi et al. (2016)
	thermosensitive male-sterile	CRISPR/Cas9	<i>TMS5</i>	Li et al. (2017)
Wheat	grain yield (seed size and thousand grain weight)	CRISPR/Cas9	<i>GW2, LPX-1, MLO</i>	Wang et al., 2018
	grain yield	CRISPR/Cas9	<i>GASR7</i>	Zhang et al. (2016)
	low gluten wheat for reduced allergenicity	CRISPR/Cas9	<i>Alpha-gliadin array, Gli-2 locus</i>	Sanchez-Leon et al. (2018)
	powdery mildew resistance	CRISPR/Cas9	<i>EDR1</i>	Zhang et al. (2017)
	grain quality in hardness, starch composition and dough colour	CRISPR/Cas9	<i>pinb, waxy, ppo and psy</i>	Zhang et al. (2021)
	increased phosphorus uptake	CRISPR/Cas9	<i>PHO2</i>	Ouyang et al. (2016)
	Increased nutritional quality (biofortification)	CRISPR/Cas9	<i>VIT</i>	Connorton et al. (2017)
	increased abiotic stress	CRISPR/Cas9	<i>DREB, ERF</i>	Kim et al. (2018)

INTERNATIONAL REGULATION

The rapid development of these new plant breeding techniques has led to the emergence of several aspects related to the state of regulation of plants obtained by genome editing.

According to many scientists, the modifications made by CRISPR are no different from natural or conventional breeding, and thus CRISPR-edited varieties should not be subject to the same regulations as GMOs (Pacher and Puchta, 2017).

However, many countries still rely on the same GMO regulations, and others are reviewing and developing new regulations for GMOs and genome editing products.

Australia, Japan and USA have decided not to regulate the genome-edited products. Brazil, Argentina and Chile have partial regulations but largely do not regulate genome editing products. Canada, Mexico, New Zealand, India, Malaysia, South Africa, Thailand and EU regulate both GMOs and genome-editing products, but here there is debate and re-evaluation to arrive at the best solution (Prasetya and Nugroho, 2021).

In July 2018, the Court of Justice of the European Union ruled that gene-edited crops must be regulated as genetically modified organisms.

CONCLUSIONS

World population increase, along with climate change, has exacerbated the problem of food shortages. Obtaining new crop varieties through conventional breeding methods can be a lengthy process that cannot provide the rapid progress needed to ensure food security and tackle climate change.

Genome editing technologies are new and innovative tools of great importance for plant breeding, the key elements of their efficiency being the precision and the significant reduction of the time to obtain the final product.

In recent years, the CRISPR/Cas9 technique has been widely used in maize and wheat breeding to develop some essential agronomic traits and to combat abiotic and biotic stress.

The success of obtaining new crop varieties through this technique will largely depend on the regulations for the use of such plants.

REFERENCES

- Bonea, D., (2013). *Plante modificate genetic. Obținere și utilizare*. Editura Universitaria, Craiova.
- Bonea, D., (2020). Grain yield and drought tolerance indices of maize hybrids, *Notulae Scientia Biologicae*, 12(2), 376-386.
- Bonea, D., (2021). Evolution and global distribution of genetically modified soybean area in the period 2014-2018. *Scientific Papers Series Management, Economic Engineering in Agriculture and Rural Development*, 21(4), 71-79.
- Bonea, D., (2022). Analysis of global trends in GM maize approvals in the period 2014-2018. *Scientific Papers Series Management, Economic Engineering in Agriculture and Rural Development*, 22(1), 53-59.
- Braatz, J., Harloff, H.J., Mascher, M., Stein, N., Himmelbach, A., Jung, C., (2017). CRISPR-Cas9 targeted mutagenesis leads to simultaneous modification of different homoeologous gene copies in polyploid oilseed rape

- (Brassica napus). *Plant Physiology*, 174(2), 935–942.
- Congressional Research Service (CRS), (2018). *Advanced Gene Editing: CRISPR-Cas9*. <https://crsreports.congress.gov/product/pdf/R/R44824/6>
- Connorton, J.M., Jones, E.R., Rodríguez-Ramiro, I., Fairweather-Tait, S., Uauy, C., Balk, J., (2017). Wheat vacuolar iron transporter TaVIT2 transports Fe and Mn and is effective for biofortification. *Plant Physiology*, (4), 174, 2434–2444.
- Constantinescu, E., Olaru, L.A., (2017). The productivity of some wheat varieties in the soil and climate conditions from South western zone of Mehedinți county. *Annals of the University of Craiova – Agriculture, Montanology, Cadastre Series*, vol XLVII, 99-105
- Constantinescu, E., Nițu (Jianu), R., Muscalu, A., (2021). Study regarding the behavior of some varieties of wheat, in ecopedological conditions, from the south-west area of Dolj County. *Annals of the University of Craiova – Agriculture, Montanology, Cadastre Series*, 51(1), 85-96.
- Dhankher, O.P., Foyer, C.H., (2018). Climate resilient crops for improving global food security and safety. *Plant Cell and Environment*, 41(5), 877–884.
- Doudna, J. A., Charpentier, E., (2014). The new frontier of genome engineering with CRISPR-Cas9. *Science*, 346: 1258096.
- Ezezi, O.C., Daar, A.S., Barber, K., Mabeya, J., Thomas, F., Deadman, J., Wang, D., Singer, P.A., (2012). Factors influencing agbiotech adoption and development in sub-Saharan Africa. *Nature Biotechnology*, 30(1), 38–40. <https://doi.org/10.1038/nbt.2088>
- FAOSTAT (FAO), (2022). *Food outlook. Biannual report on global food markets*. <https://www.fao.org/3/cb9427en/cb9427en.pdf>
- Gao, H., Gadlage, M. J., Lafitte, H. R., Lenderts, B., Yang, M., Schroder, M., Farrell, J., Snopek, K., Peterson, D., Feigenbutz, L., Jones, S., Clair, G.St., Rahe, M., Sanyour-Doyel, N., et al., (2020). Superior field performance of waxy corn engineered using CRISPR–Cas9. *Nature Biotechnology*, 38(5), 579–581. <https://doi.org/10.1038/s41587-020-0444-0>
- He, Y., Zhao, Y., (2020). Technological breakthroughs in generating transgene-free and genetically stable CRISPR-edited plants, *aBIOTECH*, 1, 88–96. <https://doi.org/10.1007/s42994-019-00013-x>
- Hsu, P.D., Lander, E.S., Zhang, F., (2014). Development and applications of CRISPR-Cas9 for genome engineering. *Cell*, 157, 1262–1278.
- Jinek, M., Chylinski, K., Fonfara, I., Hauer, M., Doudna, J.A., Charpentier, E.A., (2012). Programmable Dual-RNA — Guided DNA endonuclease in adaptive bacterial immunity. *Science*, 337 (6096), 816–821.
- Kim, H., Kim, J.S., (2014). A guide to genome engineering with programmable nucleases. *Nature Reviews Genetics*, 15, 321–334.
- Kim, D., Alptekin, B., Budak, H., (2018). CRISPR/Cas9 genome editing in wheat. *Functional and Integrative Genomics*, 18(1), 31–41. <https://doi.org/10.1007/s10142-017-0572-x>

- Li, J., Zhang, H., Si, X., Tian, Y., Chen, K., Liu, J., Chen, H., Gao, C., (2017). Generation of thermosensitive male-sterile maize by targeted knockout of the ZmTMS5 gene. *Journal of Genetics and Genomic (Yi Chuan Xue Bao)*, 44, 465-468.
- Liang, Z., Zhang, K., Chen, K., Gao, C., (2014). Targeted mutagenesis in Zea mays using TALENs and the CRISPR/Cas system. *Journal of Genetics and Genomics*, 41(2), 63–68.
- Liu, L., Gallagher, J., Arevalo, E.D., Chen, R., Skopelitis, T., Wu, Q., Bartlett, M., Jackson, D., (2021). Enhancing grain-yield-related traits by CRISPR–Cas9 promoter editing of maize CLE genes. *Nature Plants*, 7, 287–294.
- Ma, X., Liu, Y.G., (2016). CRISPR/Cas9-based multiplex genome editing in monocot and dicot plants. *Current Protocols in Molecular Biology*, 115, 31.
- Martin-Laffon, J., Kuntz, M., Ricoch, A.E., (2019). Worldwide CRISPR patent landscape shows strong geographical biases. *Nature Biotechnology*, 37, 601–621.
- Nuccio, M. L., Claeys, H., Heyndrickx, K.S., (2021). CRISPR-Cas technology in corn: a new key to unlock genetic knowledge and create novel products. *Molecular Breeding*, 41: 11. <https://doi.org/10.1007/s11032-021-01200-9>
- Ouyang, X., Hong, X., Zhao, X., Zhang, W., He, X., Ma, W., Teng, W., Tong, Y., (2016). Knock out of the PHOSPHATE 2 Gene TaPHO2-A1 improves Phosphorus uptake and grain yield under low Phosphorus conditions in common wheat. *Scientific Reports*, 6: 29850. <https://doi.org/10.1038/srep29850>
- Pacher, M., Puchta, H., (2017). From classical mutagenesis to nuclease-based breeding – directing natural DNA repair for a natural end-product. *The Plant Journal*, 90(4), 819–833.
- Pickar-Oliver, A., Gersbach, C.A., (2019). The next generation of CRISPR–Cas technologies and applications. *Nature Reviews Molecular Cell Biology*, 20(8),490–507.
- Prasetya, B., Nugroho, S., (2021). *The role of genome editing to boost bioeconomy significantly: opportunities and challenges in Indonesia*. Proceedings The 5th SATREPS Conference, 3(1),47-62.
- Qi, W., Zhu, T., Tian, Z., Li, C., Zhang, W., Song, R., (2016). High-efficiency CRISPR/Cas9 multiplex gene editing using the glycine tRNA-processing system-based strategy in maize. *BMC Biotechnology*, 16, 58. <https://doi.org/10.1186/s12896-016-0289-2>
- Rasheed, A., Gill, R.A., Hassan, M.U., Mahmood, A., Qari, S., Zaman, Q.U., Ilyas, M., Aamer, M., Batool, M., Li, H., Wu, Z.A., (2021). A critical review: Recent advancements in the use of CRISPR/Cas9 technology to enhance crops and alleviate global food crises. *Current Issues in Molecular Biology*, 43(3),1950–1976.
- Ritchie, H., Roser, M., (2019). *Land Use. 2019*. Published online at Our World In Data. org. <https://ourworldindata.org/land-use>
- Sánchez-León, S., Gil-Humanes, J., Ozuna, C.V., Giménez, M.J., Sousa, C., Voytas, D.F., Barro, F., (2018). Low-Gluten, nontransgenic wheat engineered with CRISPR/Cas9. *Plant Biotechnology*, 16, 902–910.
- Shi, J., Gao, H., Wang, H., Lafitte, H.R., Archibald, R.L., Yang, M., Hakimi, S.M., Mo, H., Habben, J.E., (2017). ARGOS 8 variants generated by CRISPR-Cas9

- improve maize grain yield under field drought stress conditions. *Plant Biotechnology Journal*, 15(2), 207–216.
- Ștefan, M., Constantinescu, E., (2011). *Fitotehnie: anii de studiu III-IV: manual universitar pentru învățământul la distanță*, Editura Universitaria, Craiova.
- Urechean, V., Bonea, D., (2017). Coexistence in cultivation of genetically modified maize (MON810) with conventional maize. *Romanian Agricultural Research*, 34, 51-58.
- Wang, W., Pan, Q., He, F., Akhunova, A., Chao, S., Trick, H., Akhunov, E., (2018). Transgenerational CRISPR-Cas9 activity facilitates multiplex gene editing in allopolyploid wheat. *The CRISPR Journal*, 1(1), 65–74.
<https://doi.org/10.1089/crispr.2017.0010>
- Zaidi, S.SeA., Mahas, A., Vanderschuren, H., Mahfouz M.M., (2020). Engineering crops of the future: CRISPR approaches to develop climate-resilient and disease-resistant plants. *Genome Biology*, 21, 289.
<https://doi.org/10.1186/s13059-020-02204-y>
- Zhang, Y., Liang, Z., Zong, Y., Wang, Y., Liu, J., Chen, K., Qiu, J., Gao, C., (2016). Efficient and transgene-free genome editing in wheat through transient expression of CRISPR/Cas9 DNA or RNA. *Nature Communications*, 7,12617.
<https://doi.org/10.1038/ncomms12617>
- Zhang, Y., Bai, Y., Wu, G., Zou, S., Chen, Y., Gao, C., Tang, D., (2017). Simultaneous modification of three homoeologs of TaEDR1 by genome editing enhances powdery mildew resistance in wheat. *The Plant Journal*, 91(4),714–724.
- Zhang, M., Cao, Y., Wang, Z., Wang, Z.Q., Shi, J., Liang, X., Song, W., Chen, Q., Lai, J., Jiang, C.A., (2018). A retrotransposon in an HKT1 family sodium transporter causes variation of leaf Na⁺ exclusion and salt tolerance in maize. *The New Phytologist*, 217(3),1161–1176.
- Zhang, S., Zhang, R., Gao, J., Song, G., Li, J., Li, W., Qi, Y., Li, Y., Li, G., (2021). CRISPR/Cas9 - mediated genome editing for wheat grain quality improvement. *Plant Biotechnology Journal*, 19, 1684–1686.