

Review On How Genetic Innovations Revolutionizing Seed Quality Production

Abstract

The process of genetic engineering modifies the genes regulating various attributes of seed quality by fabricating or redesigning some of the DNA sequences within an organism which affects germination, growth, and storage performance. Providing molecular dissection as a tool for exploiting the genetic blueprint towards the crop improvement strategy as well as improved seed yield or quality is an example showing how important it is to understand these things technically. Techniques like CRISPR and prime editing that have potential to position-specific changes in DNA sequences that helped to improve different traits such as stress resistance, productivity, nutritional value *etc.* Applications of transgenic approaches for boosting mineral content in certain crops different methods used to deal with micronutrient for malnutrition in developing countries. Development of good quality seeds resistant to environmental conditions are very important for agricultural sustainability under climate change. It ensures crop adaptation, food security and multiplication that must be promoted in future. The regulatory framework rightly requires safety assessments adapted to genetically modified (GM) foodstuffs which not only leads to satisfied public regarding health issues but also ethical standards concerning genetic engineering are upheld. The outcome is that by adherence to professional ethics and strict safety measures, biotechnology possessing such qualities has been able to make revolutionary contribution into development fostered by stable agriculture system.

Keywords: *Genetic Engineering, Seed Quality, Precision Editing, and Biofortification.*

1. Introduction

Genetic engineering plays a crucial role in enhancing seed quality by manipulating genetic factors that influence seed performance. Seed quality is defined by a combination of genetic, physical, physiological, and sanitary attributes that impact germination, vigor, and longevity (Moterleet *et al.*, 2011). Seed metabolites are directly linked to seed quality and nutritional value, making knowledge of seed chemical compositions essential for modern agricultural breeding programs (Lin *et al.*, 2014). The ability of seeds to germinate under diverse environmental conditions and develop into healthy seedlings is a key aspect of seed quality (Kazmi *et al.*, 2011). Genetic engineering provides opportunities to improve seed quality by modifying genes to enhance agronomic performance, resistance to pests and diseases, and end-use quality of crops (Shewry & Lazzeri, 1996). Through genetic manipulation, it is possible to enhance seed protein content by modulating biosynthetic pathways and improving nitrogen relocation to the grain (Wenefrida *et al.*, 2013). Additionally, genetic engineering can introduce traits like apomixis into crop plants, resulting in the production of clonal seeds identical to the mother plant (Rueda *et al.*, 2016).

The genetic makeup of seeds is intricate, involving interactions between molecular processes, seed characteristics, and environmental cues to ensure optimal germination timing for plant survival (Joosen *et al.*, 2011). The genetic program governing seed production is evident in various parts of the seed, including the embryo, endosperm, and maternal plant (Herman, 2014). Understanding the genetic diversity within crop germplasm is essential for crop improvement, enabling the conservation of best-performing cultivars and wild races, which are crucial elements in breeding programs (Shah *et al.*, 2018). So genetic engineering significantly contributes to improving seed quality by manipulating genetic factors that influence seed performance. Understanding the genetic basis of seed quality enables the development of crops with enhanced traits such as higher protein content, disease resistance, and optimal germination timing, thereby advancing agriculture and food production.

2. Understanding the Genetic Blueprint

Understanding the genetic blueprint is essential for enhancing seed quality in plants. Research has demonstrated that elucidating the genetic regulatory mechanisms involved in seed development can lead to strategies for improving yield and quality (Chen *et al.*, 2014). Molecular approaches, such as marker-assisted selection and 'omics' research, play a significant role in enhancing seed quality parameters in crops like soybeans (Tripathi & Khare, 2016). Knowledge of the cultivar effect on seed oil quality is crucial for improving oilseed plants genetically (Kaseke *et al.*, 2020). By unraveling the complex traits of seed quality through genetics, physiology, and -omics technologies, it is possible to identify regulatory genes and pathways to predict and enhance seed quality (Ligterink *et al.*, 2012).

Seed dormancy is a desirable trait in crops, and understanding the genetic and molecular mechanisms involved is crucial for improving seed quality in breeding programs (Zhang *et al.*, 2020). Research focusing on seed germination and vigor highlights the importance of selecting quality seeds, priming seeds, and breeding varieties with improved seed performance to ensure crop sustainability (Reed *et al.*, 2022). Spatiotemporal transcriptomics and metabolic profiling provide insights into gene regulatory networks during seed development, which is crucial for genetic improvement of crop yield and quality (Yu *et al.*, 2023). Studies have shown that seed mass increases with higher pollen loads, indicating enhanced seed quality through increased pollen competition or genetic sampling (Hildesheim *et al.*, 2019). Identifying genetic markers for increased protein content in seeds can lead to the development of new plant varieties with improved nutritional value (Grimberg *et al.*, 2022). The molecular basis of seed longevity, including candidate genes and proteins associated with seed longevity, is essential for understanding seed quality (Ramtekey *et al.*, 2022). Seed endophytes play a vital role in plant breeding strategies, as seeds carry the genetic blueprint of plants and harbor diverse microbiota (Gopal & Gupta, 2016).

3. Precision Editing

Precision editing techniques such as prime editing and CRISPR have transformed genome editing, allowing for precise modifications in DNA sequences (Anzalone *et al.*, 2019; Gaba *et al.*, 2021). These methods facilitate the direct incorporation of new genetic information into specific DNA sites, enabling targeted enhancements in crop traits (Sedeeke *et al.*, 2019; Li *et*

et al., 2020). Through the use of prime editing, gene replacement can be achieved without double-strand breaks, enhancing the potential for precise targeted gene/allele replacement in crops (Li *et al.*, 2020). These advancements offer promise for boosting crop productivity and resilience to climate change (Sedeek *et al.*, 2019).

Genome editing tools like CRISPR/Cas have demonstrated significant potential in enhancing abiotic stress resilience in plants (Kaur *et al.*, 2022). These tools present opportunities to alter plants for improved stress tolerance, essential for sustainable agriculture in response to changing environmental conditions (Kaur *et al.*, 2022). Additionally, the application of genome editing technologies has paved the way for identifying pathways for desirable trait modification, underscoring the significance of accurately integrating valuable traits and eliminating undesirable ones (Picheny *et al.*, 2017; Gaba *et al.*, 2021).

Moreover, the implementation of genome editing in agriculture has resulted in the development of plants with enhanced food and feed quality, increased yield, and tolerance to biotic and abiotic stress (Modrzejewski *et al.*, 2019). These advancements underscore the potential of genome editing as a tool for precise plant trait modification, offering novel avenues for improving crop traits and overall agricultural resilience (Modrzejewski *et al.*, 2019). So the precision editing techniques like prime editing and CRISPR have introduced new possibilities for tailored trait modifications in crops, providing precise and effective methods for enhancing performance and resilience in agricultural settings.

4. Biofortification

Biofortification is a strategy that aims to enhance the nutritional quality of staple crops through genetic manipulation, including conventional breeding and transgenic approaches (Yadava *et al.*, 2018). This approach involves increasing the concentration of target nutrients in food crops to address prevalent nutrient deficiencies, particularly in regions where malnutrition is a significant concern (Siwela *et al.*, 2020). By harnessing the powers of plant breeding, biofortification seeks to improve the bioavailability of essential nutrients in edible plant parts, making it a preferred method for enriching grains with vitamins and minerals (Liberal *et al.*, 2020).

Through genetic manipulation, biofortification can increase the micronutrient density of staple crops, offering a promising solution to combat hidden hunger and improve the health of populations in both rural and urban areas of the developing world (Pfeiffer & McClafferty, 2007). Studies have shown that genetic engineering can significantly enhance the nutritional qualities of plants, surpassing the results achieved through conventional breeding methods (Naqvi *et al.*, 2009). By increasing the mineral nutrient content in crops, biofortification can be achieved through agronomic practices, such as fertilization, or genetic manipulation to develop cultivars with higher mineral absorption capabilities (Shoormijet *et al.*, 2022).

Biofortification strategies involve crop breeding, genetic manipulation, and the application of mineral fertilizers to address micronutrient malnourishment effectively (Chaudhari *et al.*, 2022). Genetic biofortification combines conventional and transgenic breeding techniques to introgress genes that promote high micronutrient accumulation into elite genotypes or manipulate crop genetic makeup using genome editing tools to enhance nutrient levels (Tumbare & Maphosa,

2023). The ultimate goal of biofortification is to elevate mineral content, improve fatty acid composition, increase amino acid levels, and enhance antioxidant levels in food crops (Hirschi, 2009).

5. Environmental Adaptation

To enhance sustainability in changing climates, it is crucial to focus on engineering seeds that exhibit climate resilience. Climate resilience in crops involves both short-term management by seed traders and farmers through cultivar selection and long-term contributions by breeders in providing diverse cultivar responses (Kahiluoto *et al.*, 2018). Strategies such as utilizing improved or hybrid seeds and exchanging seeds among villages have been identified as effective climate-resilience approaches for small-scale producers in low- and middle-income countries (Acevedo *et al.*, 2020). Moreover, the development of seeds resilient to current and future climate shocks is highlighted as a critical adaptation option for smallholder farmers in sub-Saharan Africa (Cacho *et al.*, 2020).

Climate change significantly impacts plant regeneration from seeds by altering environmental cues, which can affect the timing and success of seed germination and growth (Walck *et al.*, 2011). Understanding the response and adaptive capacity of seed banks to climate change is essential for predicting future species distributions and risks of extinction in ecosystems subjected to disturbances (Ooi, 2012). Additionally, integrating genebanks with farmers through the distribution of climate-resilient seed kits can aid in promoting crop variety adoption and understanding the effectiveness of such interventions (Stoilova *et al.*, 2019). Efforts to build resilience in food systems and enhance societal and ecological resilience through plant breeding are emphasized (Bueren *et al.*, 2018). Breeders are challenged to produce cultivars that strengthen ecological and societal resilience by aiming for sustainability targets such as food security, agrobiodiversity, and climate robustness (Bueren *et al.*, 2018). By focusing on climate-resilient seeds and promoting seed diversity, it is possible to improve agricultural sustainability and adaptability to changing climates.

6. Regulatory Landscape

Genetic engineering involves a regulatory framework that mandates safety assessments and consultations with multiple regulatory agencies before genetically engineered (GE) crops can be commercialized and grown in the US (Hood *et al.*, 2019). This stringent process highlights the emphasis placed on ensuring the safety of genetically modified organisms and the environment. Public concerns surrounding genetic engineering encompass various issues related to risk, benefit, and ethics, underscoring the importance of understanding the public discourse on this technology (Frewer *et al.*, 1997). Ethical considerations related to genetic engineering have sparked debates ranging from the practical applications of genetic engineering to the ethical implications of direct-to-consumer genetic testing (Allyse *et al.*, 2018; Mandal & Bach, 2018).

The ethical framework of genetic engineering is based on considerations of risk, benefit, and broader societal implications. While some argue that genetic engineering itself may not pose intrinsic ethical issues, concerns about potential misuse raise ethical considerations that require careful attention (Lucassen, 1996). The ethical discourse on genetic engineering extends beyond

scientific realms to intersect with theological perspectives, as evidenced by the evaluation of genetic engineering through the lens of bioethics (Watling, 2006).

Education plays a vital role in shaping ethical awareness within the field of genetic engineering. Teaching the ethics of genetic engineering is crucial for promoting a responsible approach to the development and application of genetic technologies (Lucassen, 1995). Moreover, there have been calls for the establishment of a universal code of ethics for medical and biological engineers to uphold ethical standards in genetic engineering practices (Voigt, 2010). That's why a comprehensive understanding of the regulatory landscape, public concerns, ethical frameworks, and the role of education is essential for navigating the ethical and legal dimensions of genetic engineering responsibly.

Conclusion

The genetic engineering stands as a pivotal tool in enhancing seed quality, with the understanding of genetic, physical, physiological, and sanitary attributes that influence seed performance. Now a days manipulation in genetics, agronomic traits, resistance to pests and diseases, and nutritional value of crops is very easy by genetic engineering, thus addressing critical agricultural needs. The complexity of the genetic blueprint governing seed development necessitates advanced molecular approaches, such as marker-assisted selection and 'omics' technologies, to elucidate regulatory mechanisms and improve seed traits like dormancy, germination, and protein content. Precision editing techniques, notably CRISPR and prime editing, have revolutionized genome editing by enabling precise modifications in plant DNA. These advancements hold immense potential for enhancing crop productivity and resilience, particularly in the face of climate change. Genome editing also helps to biofortification and modification in gene combination of crop plants that can lead to development of new desired crops with useful traits, it offers a viable strategy to combat malnutrition by increasing the nutrient density of staple crops. These approaches significantly improve the health of population in regions or countries where nutrient deficiencies are prevalent. Biofortification enhances the micronutrient content in crops, thus providing a sustainable solution to hidden hunger and contributing to food security. Environmental adaptation strategies focus on developing seeds that can withstand climate variability, ensuring sustainability in agriculture. This involves breeding cultivars with diverse responses to environmental stresses and promoting seed diversity. The integration of genebanks and the distribution of climate-resilient seed kits are vital for enhancing the adaptability of crops to changing climates, particularly for smallholder farmers in vulnerable regions. The regulatory landscape of genetic engineering is stringent, requiring safety assessments and consultations to ensure the safe commercialization of GE crops.

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References:

- Acevedo, M., Pixley, K., Zinyengere, N., Meng, S., Tufan, H., Cichy, K., ... & Porciello, J. (2020). A scoping review of adoption of climate-resilient crops by small-scale producers in low- and middle-income countries. *Nature Plants*, 6(10), 1231-1241. <https://doi.org/10.1038/s41477-020-00783-z>
- Allyse, M., Robinson, D., Ferber, M., & Sharp, R. (2018). Direct-to-consumer testing 2.0: emerging models off direct-to-consumer genetic testing. *Mayo Clinic Proceedings*, 93(1), 113-120. <https://doi.org/10.1016/j.mayocp.2017.11.001>
- Anzalone, A., Randolph, P., Davis, J., Sousa, A., Koblan, L., Levy, J., ... & Liu, D. (2019). Search-and-replace genome editing without double-strand breaks or donor dna. *Nature*, 576(7785), 149-157. <https://doi.org/10.1038/s41586-019-1711-4>
- Bueren, E., Struik, P., Eekeren, N., & Nuijten, E. (2018). Towards resilience through systems-based plant breeding. a review. *Agronomy for Sustainable Development*, 38(5). <https://doi.org/10.1007/s13593-018-0522-6>
- Cacho, O., Moss, J., Thornton, P., Herrero, M., Henderson, B., BDIRSKY, B., ... & Lipper, L. (2020). The value of climate-resilient seeds for smallholder adaptation in sub-saharanafrica. *Climatic Change*, 162(3), 1213-1229. <https://doi.org/10.1007/s10584-020-02817-z>
- Chaudhari, G., Patel, D., Parmar, D., & Patel, K. (2022). Fe, zn&amp; protein content in grain, per se performance, heterosis, combining ability of grain yield in bread wheat (*triticum aestivum*) under normal &amp; late sowing condition... <https://doi.org/10.21203/rs.3.rs-1338914/v1>
- Chen, J., Zeng, B., Zhang, M., Xie, S., Wang, G., Hauck, A., ... & Lai, J. (2014). Dynamic transcriptome landscape of maize embryo and endosperm development. *Plant Physiology*, 166(1), 252-264. <https://doi.org/10.1104/pp.114.240689>
- Frewer, L., Howard, C., & Shepherd, R. (1997). Public concerns in the united kingdom about general and specific applications of genetic engineering: risk, benefit, and ethics. *Science*

- Gaba, Y., Pareek, A., & Sneh, L. (2021). Raising climate-resilient crops: journey from the conventional breeding
- Gopal, M. and Gupta, A. (2016). Microbiome selection could spur next-generation plant breeding strategies. *Frontiers in Microbiology*, 7. <https://doi.org/10.3389/fmicb.2016.01971>
- Grimberg, Å., Saripella, G., Valencia, R., Bengtsson, T., Alandia, G., & Carlsson, A. (2022). Transcriptional regulation of quinoa seed quality: identification of novel candidate genetic markers for increased protein content. *Frontiers in Plant Science*, 13. <https://doi.org/10.3389/fpls.2022.816425>
- Herman, E. (2014). Soybean seed proteome rebalancing. *Frontiers in Plant Science*, 5. <https://doi.org/10.3389/fpls.2014.00437>
- Hildesheim, L., Opedal, Ø., Armbruster, W., & Pélabon, C. (2019). Quantitative and qualitative consequences of reduced pollen loads in a mixed mating plant. *Ecology and Evolution*, 9(24), 14253-14260. <https://doi.org/10.1002/ece3.5858>
- Hirschi, K. (2009). Nutrient biofortification of food crops. *Annual Review of Nutrition*, 29(1), 401-421. <https://doi.org/10.1146/annurev-nutr-080508-141143>
- Hood, E., Eversole, K., Leach, L., Hogan, M., McHughen, A., Cordts, J., ... & Ireby, M. (2019). Report on the scra nuts and bolts workshop ii: case studies of citrus greening, ultra-low gossypol cotton, and blight tolerant, low-acrylamide potato. *Gm Crops & Food*, 10(3), 139-158. <https://doi.org/10.1080/21645698.2019.1634989>
- Joosen, R., Arends, D., Willems, L., Ligterink, W., Jansen, R., & Hilhorst, H. (2011). Visualizing the genetic landscape of arabidopsis seed performance. *Plant Physiology*, 158(2), 570-589. <https://doi.org/10.1104/pp.111.186676>
- Kahiluoto, H., Kaseva, J., Bálék, J., Olesen, J., Ruiz-Ramos, M., Gobin, A., ... & Trnka, M. (2018). Decline in climate resilience of european wheat. *Proceedings of the National Academy of Sciences*, 116(1), 123-128. <https://doi.org/10.1073/pnas.1804387115>
- Kaseke, T., Opara, U., & Fawole, O. (2020). Fatty acid composition, bioactive phytochemicals, antioxidant properties and oxidative stability of edible fruit seed oil: effect of preharvest and processing factors. *Heliyon*, 6(9), e04962. <https://doi.org/10.1016/j.heliyon.2020.e04962>
- Kaur, N., Sharma, S., Hasanuzzaman, M., & Pati, P. (2022). Genome editing: a promising approach for achieving abiotic stress tolerance in plants. *International Journal of Genomics*, 2022, 1-12. <https://doi.org/10.1155/2022/5547231>
- Kazmi, R., Khan, N., Willems, L., Heusden, A., Ligterink, W., & Hilhorst, H. (2011). Complex genetics controls natural variation among seed quality phenotypes in a recombinant inbred population of an interspecific cross between *solanum lycopersicum* × *solanum*

- pimpinellifolium. *Plant Cell & Environment*, 35(5), 929-951. <https://doi.org/10.1111/j.1365-3040.2011.02463.x>
- Li, J., Li, H., Chen, J., Yan, L., & Xia, L. (2020). Toward precision genome editing in crop plants. *Molecular Plant*, 13(6), 811-813. <https://doi.org/10.1016/j.molp.2020.04.008>
- Liberal, Â., Pinela, J., Quintana, A., Ferreira, I., & Barros, L. (2020). Fighting iron-deficiency anemia: innovations in food fortificants and biofortification strategies. *Foods*, 9(12), 1871. <https://doi.org/10.3390/foods9121871>
- Ligterink, W., Joosen, R., & Hilhorst, H. (2012). Unravelling the complex trait of seed quality: using natural variation through a combination of physiology, genetics and -omics technologies. *Seed Science Research*, 22(S1), S45-S52. <https://doi.org/10.1017/s0960258511000328>
- Lin, H., Rao, J., Shi, J., Hu, C., Cheng, F., Wilson, Z., ... & Quan, S. (2014). Seed metabolomic study reveals significant metabolite variations and correlations among different soybean cultivars. *Journal of Integrative Plant Biology*, 56(9), 826-836. <https://doi.org/10.1111/jipb.12228>
- Lucassen, E. (1995). Teaching the ethics of genetic engineering. *Journal of Biological Education*, 29(2), 129-138. <https://doi.org/10.1080/00219266.1995.9655432>
- Lucassen, E. (1996). The ethics of genetic engineering. *Journal of Applied Philosophy*, 13(1), 51-62. <https://doi.org/10.1111/j.1468-5930.1996.tb00149.x>
- Mandal, S. and Bach, C. (2018). Genetic engineering -tool for mankind. *European Journal of Engineering and Technology Research*, 3(3), 58. <https://doi.org/10.24018/ejers.2018.3.3.551>
- Modrzejewski, D., Hartung, F., Sprink, T., Krause, D., Kohl, C., & Wilhelm, R. (2019). What is the available evidence for the range of applications of genome-editing as a new tool for plant trait modification and the potential occurrence of associated off-target effects: a systematic map. *Environmental Evidence*, 8(1). <https://doi.org/10.1186/s13750-019-0171-5>
- Moterle, L., Braccini, A., Scapim, C., Pinto, R., Gonçalves, L., Rodrigues, R., ... & Júnior, A. (2011). Combining ability of popcorn lines for seed quality and agronomic traits. *Euphytica*, 185(3), 337-347. <https://doi.org/10.1007/s10681-011-0458-2>
- Naqvi, S., Zhu, C., Farré, G., Ramessar, K., Bassié, L., Breitenbach, J., ... & Christou, P. (2009). Transgenic multivitamin corn through biofortification of endosperm with three vitamins representing three distinct metabolic pathways. *Proceedings of the National Academy of Sciences*, 106(19), 7762-7767. <https://doi.org/10.1073/pnas.0901412106>
- Ooi, M. (2012). Seed bank persistence and climate change. *Seed Science Research*, 22(S1), S53-S60. <https://doi.org/10.1017/s0960258511000407>
- Pfeiffer, W. and McClafferty, B. (2007). Harvestplus: breeding crops for better nutrition. *Crop Science*, 47(S3). <https://doi.org/10.2135/cropsci2007.09.0020ipbs>

- Picheny, V., Casadebaig, P., Trépos, R., Faivre, R., Silva, D., Vincourt, P., ... & Costes, E. (2017). Using numerical plant models and phenotypic correlation space to design achievable ideotypes. *Plant Cell & Environment*, 40(9), 1926-1939. <https://doi.org/10.1111/pce.13001>
- Ramtekey, V., Cherukuri, S., Kumar, S., V. S., Sheoran, S., K., U., ... & Singh, H. (2022). Seed longevity in legumes: deeper insights into mechanisms and molecular perspectives. *Frontiers in Plant Science*, 13. <https://doi.org/10.3389/fpls.2022.918206>
- Reed, R., Bradford, K., & Khanday, I. (2022). Seed germination and vigor: ensuring crop sustainability in a changing climate. *Heredity*, 128(6), 450-459. <https://doi.org/10.1038/s41437-022-00497-2>
- Rueda, A., Grossniklaus, U., & Schmidt, A. (2016). Laser-assisted microdissection (lam) as a tool for transcriptional profiling of individual cell types. *Journal of Visualized Experiments*, (111). <https://doi.org/10.3791/53916>
- Sedeek, K., Mahas, A., & Mahfouz, M. (2019). Plant genome engineering for targeted improvement of crop traits. *Frontiers in Plant Science*, 10. <https://doi.org/10.3389/fpls.2019.00114>
- Shah, S., Khan, S., Ullah, K., & Sayal, O. (2018). Genetic diversity in cotton germplasm using multivariate analysis. *Sarhad Journal of Agriculture*, 43(1). <https://doi.org/10.17582/journal.sja/2018/34.1.130.135>
- Shewry, P. and Lazzeri, P. (1996). Genetic manipulation of crops. *British Food Journal*, 98(4/5), 5-9. <https://doi.org/10.1108/00070709610119801>
- Shoormij, F., Mirlohi, A., Saeidi, G., & Shirvani, M. (2022). Combined foliar application of zn and fe increases grain micronutrient concentrations and alleviates water stress across diverse wheat species and ploidal levels. *Scientific Reports*, 12(1). <https://doi.org/10.1038/s41598-022-24868-1>
- Siwela, M., Pillay, K., Govender, L., Lottering, S., Mudau, F., Modi, A., ... & Mabhaudhi, T. (2020). Biofortified crops for combating hidden hunger in south africa: availability, acceptability, micronutrient retention and bioavailability. *Foods*, 9(6), 815. <https://doi.org/10.3390/foods9060815>
- Stoilova, T., Zonneveld, M., Roothaert, R., & Schreinemachers, P. (2019). Connecting genebanks to farmers in east africa through the distribution of vegetable seed kits. *Plant Genetic Resources*, 17(03), 306-309. <https://doi.org/10.1017/s1479262119000017>
- to new breeding approaches. *Current Genomics*, 22(6), 450-467. <https://doi.org/10.2174/1389202922666210928151247>
- Tripathi, N. and Khare, D. (2016). Molecular approaches for genetic improvement of seed quality and characterization of genetic diversity in soybean: a critical review. *Biotechnology Letters*, 38(10), 1645-1654. <https://doi.org/10.1007/s10529-016-2154-8>
- Tumbare, T. and Maphosa, M. (2023). Genetic bio-fortification of cereals from a plant breeding perspective. *Journal of Plant Breeding and Crop Science*, 15(1), 42-56. <https://doi.org/10.5897/jpbcs2022.0997>

- Voigt, H. (2010). Editorial: a need for a universal code of ethics for medical and biological engineers. *Ethics in Biology Engineering and Medicine an International Journal*, 1(2), 79-81. <https://doi.org/10.1615/ethicsbiologyengmed.v1.i2.10>
- Walck, J., Hidayati, S., Dixon, K., Thompson, K., & Poschlod, P. (2011). Climate change and plant regeneration from seed. *Global Change Biology*, 17(6), 2145-2161. <https://doi.org/10.1111/j.1365-2486.2010.02368.x>
- Wenefrida, I., Utomo, H., & Linscombe, S. (2013). Mutational breeding and genetic engineering in the development of high grain protein content. *Journal of Agricultural and Food Chemistry*, 61(48), 11702-11710. <https://doi.org/10.1021/jf4016812>
- Yadava, D., Hossain, F., & Mohapatra, T. (2018). Nutritional security through crop biofortification in india: status & future prospects. *The Indian Journal of Medical Research*, 148(5), 621. https://doi.org/10.4103/ijmr.ijmr_1893_18
- Yu, B., Gao, P., Song, J., Yang, H., Qin, L., Song, H., ... & Xiang, D. (2023). Spatiotemporal transcriptomics and metabolic profiling provide insights into gene regulatory networks during lentil seed development. *The Plant Journal*, 115(1), 253-274. <https://doi.org/10.1111/tpj.16205>
- Zhang, C., Yuan, Z., Wang, Y., Sun, W., Tang, X., Sun, Y., ... & Yu, S. (2020). Genetic dissection of seed dormancy in rice (*oryza sativa* L.) by using two mapping populations derived from common parents. *Rice*, 13(1). <https://doi.org/10.1186/s12284-020-00413-4>