

REVIEW

 OPEN ACCESS



Challenges and prospects in using biotechnological interventions in *O. glaberrima*, an African cultivated rice

Gideon Sadikiel Mmbando 

Department of Biology, College of Natural and Mathematical Sciences, University of Dodoma (Udom), Dodoma, Tanzania

ABSTRACT

Africa has the world's fastest rate of population expansion, making it vulnerable to food shortages. Africa cultivates two types of rice (Asian rice; *Oryza sativa* and African rice; *Oryza glaberrima*). Native African rice called *O. glaberrima* has some intriguing characteristics, including resistance to several biotic and abiotic regional restrictions in Africa. However, *O. glaberrima* is solely employed as a tool to increase the production of *O. sativa*, which cannot grow in Africa, due to its low yield, lodging, grain breaking, and poor tissue culture ability. Enhancing breeding efforts for *O. glaberrima* is therefore critically important. The protocols for transformation and regeneration, however, are mostly for *O. sativa* and not *O. glaberrima*. This study examines the present problems with transformation and regeneration for African rice species as well as potential solutions for using modern breeding methods in *O. glaberrima*.

ARTICLE HISTORY

Received 11 Sep 2022
Revised 09 Nov 2022
Accepted 10 Nov 2022

KEYWORDS

Gene editing; genetic engineering; *O. glaberrima*; regeneration; transformation

Introduction

The phrase “Rice is life” captures the importance of this ancient grain to people as a main source of nutrition as well as spiritual and cultural nourishment, not just in Asia¹ but also in Africa. More than half of the world's population is fed by rice, making it the most significant cereal crop.² In addition, rice has developed into a model monocot system for genetic and functional genomic research.³ Only two of the 23 species in the genus *Oryza* have been grown for food: *Oryza sativa* and *Oryza glaberrima*.⁴ *O. glaberrima* Steud. was domesticated in West Africa,⁵ whereas *O. sativa* was in Asia.⁶ Both species are grown in Africa, and varieties of *O. sativa* were brought to West Africa as a result of trade between Africa and India.⁷ *O. glaberrima*'s yield is thought to be lower than *O. sativa*'s, probably as a result of many features such as lodging and grain breaking,⁸ and as a result, *O. sativa* has continued to take over its growing regions.⁸ *O. glaberrima*, however, is still grown in Africa today⁹ and is preferred by local farmers over *O. sativa* for some reasons, including resilience to a variety of regional restrictions, taste, hardness, and resistance to many biotic and abiotic stresses.¹⁰ It's been reported that farmers abandoned better

O. sativa cultivars to resume the cultivation of *O. glaberrima*. For instance, *O. glaberrima* is currently grown in a deep water location at Banfora in Burkina Faso, despite the government having already introduced a deep water variety from Indonesia¹¹; demonstrating the strong resilience of *O. glaberrima* to such severe environments, including deep water areas. However, there is a lack of an appropriate breeding and genetic development program to benefit from *O. glaberrima*'s advantages. This has kept the unwanted traits from being removed from this species while allowing the beneficial traits to be transferred to *O. sativa*, which is not an African native and cannot survive there.

Although it has been reported that the New Rice for Africa (NERICA) varieties combine some ability to develop well in the challenging environment of *O. glaberrima* as well as the high-yielding potential of *O. sativa*^{5,12,13}; they still lack resistance mechanisms to some local constraints, including weeds, in comparison to *O. glaberrima* variety.¹⁴ As a result, Africa Rice continues to look for *O. glaberrima* lines for crucial characteristics in rainfed rice in Sub-Saharan Africa (SSA). *O. glaberrima* has, however, received far less

CONTACT Gideon Sadikiel Mmbando  gideonmmbando@gmail.com  Department of Biology, College of Natural and Mathematical Sciences, University of Dodoma (Udom), P. O. BOX 259 Dodoma, Tanzania

© 2022 The Author(s). Published with license by Taylor & Francis Group, LLC.

This is an Open Access article distributed under the terms of the Creative Commons Attribution License (<http://creativecommons.org/licenses/by/4.0/>), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

scientific attention, particularly in the areas of genetic breeding and biotechnology.¹⁰ *O. glaberrima*'s undesirable traits, such as grain breaking, lodging,^{5,15,16} high ultraviolet-B (UVB) sensitivity,¹⁷ and limited yield potential, may have led to its usage primarily as a genetic resource to enhance *O. sativa*.¹⁸ To fully utilize current breeding techniques in enhancing the yield and characteristics of this species for the people of Africa and the rest of the world, it is crucial to produce appropriate transformation and regeneration of *O. glaberrima* local varieties.

This review covers the current status, challenges, and potential solutions to promote transformation and regeneration with an emphasis on African rice. This study urges the creation of a transformation and regeneration protocol for *O. glaberrima* immediately. This study also made several suggestions for how the most cutting-edge genome editing techniques, such as clustered regularly interspaced short palindromic repeats (CRISPR)-associated endonuclease Cas9 (CRISPR/Cas9), might be used to improve *O. glaberrima*'s yield and eliminate its undesirable characteristics. This is because *O. glaberrima* is an elite line that has adapted to the African environment. Such knowledge is crucial for the biotechnological advancement of African rice, which will hasten the continent's green revolution by supplying dream rice with all the nutritional benefits that developing nations most want.

1. Importance of *O. Glaberrima* to Africa

Due to climate fluctuation, rice farming in Africa is unstable and susceptible to numerous environmental challenges, including drought and submergence.¹⁹ Therefore, weak features in native and local rice cultivars that are already acclimated to other pressures in SSA should be genetically improved to create efficiently stable and sustainable agricultural systems in Africa. *O. glaberrima* is thought to have evolved tolerance to the majority of abiotic and biotic stressors for rice farming in SSA^{5,20} since it has thrived in Africa with little intervention from humans. *O. glaberrima*, for instance, is recognized as a source of resistance to a variety of native abiotic challenges, including low phosphorus availability in acid soils,²¹ submergence,^{22,23} drought,^{5,20,24,25} iron toxicity,²⁶ and biotic challenges

like nematodes,^{27–30} African rice gall midges,^{31,32} weeds,^{33–35} and the rice yellow mottle virus^{36,37} *O. glaberrima* is employed in traditional ritual rites to satisfy the souls of the ancestors, as in the communities of the Danyi plateau in Togo and the Casamance region in southern Senegal.³⁸ This is in addition to its resistance to many geographical restrictions in Africa. The genetic diversity of this species may also be an important breeding resource. For instance, increasing the amount of certain micronutrients^{10,39–41} and increasing the amount of protein in grains.^{42–44} As a result, genetically enhancing *O. glaberrima*'s poor features will have the added benefit of all these advantageous traits acting as adaptation mechanisms, which will aid the organism's ability to thrive in the stress-prone SSA region.

O. glaberrima has a great degree of adaptation to local farming in Africa, which may explain why it is still grown in some regions of West Africa today.⁹ It is a big surprise that rice researchers have neglected *O. glaberrima* breeding initiatives for many years in favor of focusing on raising the productivity of Asian rice varieties to increase food supply in both Asia and Africa, despite all the advantages described above. Very few studies have attempted to increase *O. glaberrima* yield by fusing the positive characteristics of *O. glaberrima* and *O. sativa*, as demonstrated in NERICA.^{5,12,13} What is surprising is that *O. glaberrima* has been overlooked even by African researchers studying rice and primarily uses it as a source to raise the yield of *O. sativa*. The lack and poor regeneration capacity of this species in comparison to *O. sativa* could be one explanation for this^{45,46} due to the paucity of data on *O. glaberrima*'s callus induction and plantlet regeneration.⁴⁷ The lack of suitable regeneration media may be the reason why the majority of rice researchers have given up on breeding and genetic enhancement programs for *O. glaberrima*, in addition to the sterility barrier between the two domesticated species.^{48,49} Similar to what has already been done on Asian rice varieties, rice researchers urgently need to create the ideal conditions for transformation and regeneration media for native African rice varieties. Future research should investigate this topic. The recent development in gene editing and modification allows us to artificially disrupt a single gene responsible for the reproductive barrier,^{50,51} allowing us to avoid the hybrid

sterility of *O. glaberrima* and *O. sativa* and thereby facilitate interspecific hybridization for breeding programs of the two species.

Alternatively, *O. glaberrima*'s unfavorable traits may have led to its use primarily as a genetic resource to enhance *O. sativa*.⁸ This shouldn't be a problem today, though, as recent developments in genome editing technologies, like the CRISPR/Cas 9 system,^{52–54} may make it possible to eliminate undesirable traits in *O. glaberrima* and even hasten the process of domestication.⁵⁵ The best way to improve *O. glaberrima*, for instance, maybe to use CRISPR/Cas9 technology to edit and modify the genes responsible for traits like grain shattering and lodging resistance. These traits are crucial in the improvement of *O. glaberrima*.⁵⁶ A different approach is to look for *O. glaberrima* lines that are resilient to grain lodging and shattering. Although lines that are resistant to lodging have not yet been identified,¹⁰ this has already begun. It is significant to note that proper transformation and regeneration protocols, which are thought to be crucial steps during the generation of transgenic plants,⁵⁷ are required for the success of these gene editing technologies, including CRISPR/Cas9.

2. Challenges in Improving *O. Glaberrima* through Hybridization

Genetic advancements in cereals, particularly those of economically significant crops like rice, are considered to be a promising solution to the issue of a food shortage.⁵⁸ However, the establishment of good callus induction and subsequent plant regeneration are the key steps^{59,60} before utilizing any genetic improvement program. It is well known that the accessibility of gene transfer, which is inappropriate for plant regenerations, makes it difficult for many transgenic species to recover.⁵⁷ During somatic embryogenesis, cell division and differentiation are significantly influenced by plant growth regulators.^{61,62} Therefore, embryogenic callus is essential for successful regeneration, and *O. glaberrima* may have less regeneration potential than *O. sativa* because it forms fewer embryogenic calluses. Even under different hormonal concentrations, Mmbando et al. showed that MS media⁶³

rather than N6⁶⁴ was the most effective for inducing calluses in the majority of *O. glaberrima* varieties. On the other hand, the control Asian rice cultivar Nipponbare appears to react strongly to N6 media. The variation in callus induction between Nipponbare and African rice varieties is brought on by differences in the nutrient media composition and genotype, which is one of the major factors for tissue culture.⁶⁵ Future research should test various hormonal concentration gradients using the best MS media for *O. glaberrima* suggested for cultivar TOG12380.⁶⁶

The majority of cultivars of *O. glaberrima* also showed cell necrosis, though the cause is still unknown, which prevents effective regeneration.⁶⁶ This report resembles one that was done previously on *indica* rice (*O. sativa*)⁶⁷ Numerous plant species that struggle with *in vitro* culture have been found to exhibit callus browning, which reduced their ability to grow and regenerate and even caused their death.⁶⁸ Similar to this, Mmbando et al. showed that only 1 of the 5 *O. glaberrima* tested cultivars was regenerated by Murashige and Skoog (MS)⁶¹ using media from Brisibe et al.⁶³ while the others formed tiny brown calli during the callus induction stage.⁶⁶ Maltose has been shown to cause less browning than sucrose when added to culture media.⁶⁹ Although there are numerous protocols for embryogenic calli-mediated plant regeneration in *indica* rice among *O. sativa* varieties, the majority of them are genotype-dependent, and the universal medium that can be adapted to a wide range of *indica* rice genotypes has not yet been developed.⁷⁰ The majority of callus induction protocols created for *O. sativa*, unless modified do not function well in *O. glaberrima* and frequently cause the callus to brown, necrosis, and grow poorly, all of which ultimately harm callus regeneration. Despite *O. glaberrima* tissue culture can be produced and increased in media from *O. sativa* by adjusting nutrients and growth regulators. The induction, proliferation, and regeneration of calluses on *O. sativa* are laborious and time-consuming.⁷⁰ As a result, there is a great need to create a universal medium that can be used by many different varieties of *O. glaberrima*.

3. Plant Regeneration: A Constraint in Using Biotechnological Tools for Improving *O. Glaberrima*

The smallest genome of all cereals,⁷¹ rice is the first fully sequenced agriculturally relevant crop. Additionally, *O. glaberrima*'s genome assembly and annotation have already been reported,⁷² opening the door to the possibility of fully utilizing several significant genes in this species. Although nucleus-cytoplasm interactions and various nuclear gene interactions cause a sterility barrier to exist between *O. sativa* and *O. glaberrima*,⁴⁸ recombinant DNA technology has been used as an alternative method to conventional breeding to transfer novel genes across taxonomic boundaries.⁷³ Working with living cells and their molecules is essential to biotechnology, and this can be accomplished using a variety of techniques, including molecular breeding, marker-assisted selection, mutation breeding, tissue culture, micropropagation, gene editing, and molecular diagnostic tools.⁷⁴ Since genetic engineering is the most effective method for improving rice and can introduce useful genes into rice with little disruption to the rice genome,⁷⁵ it will be the main topic of this review, along with some aspects of gene editing.

Despite the strong opposition from the anti-biotechnology lobby in Africa, local farmers have already benefited from tissue culture technologies for crops like pyrethrum, sugar cane, cassava, and bananas.⁷⁶ African rice cultivars known as NERICAs were created using embryo rescue techniques and interspecific crosses of *O. sativa* and *O. glaberrima*.⁷⁷ *O. sativa* gene transformation has been regularly carried out in some labs, though it has been indicated that the *indica* rice system is more complicated.⁷⁸ The introduction of foreign genes into plant cells and the subsequent regeneration of the transformed cells are two essential steps in the transformation of plants. The accessibility of gene transfer, which is unsuitable for plant regeneration, has made it difficult to recover transgenic species of many species.⁵⁷ Model varieties like Kasalath (*indica*) and Nipponbare (*japonica*) have established efficient rice transformation protocols using mature as well as immature seeds^{79,80}; however, the difficulties associated with sterilization and isolation of immature embryos continue to be

a serious barrier for use of these explants in transformation.⁸¹ In addition, a lot of popular food-producing varieties, like Koshihikari in Japan and IR64 in tropical regions (as well as Suakoko in Nigeria), have been shown to have lower regeneration capacity in mature seed culture systems, making it difficult to effectively improve them through genetic modification.^{82,83} Afolabi et al. were able to develop a regeneration protocol for callus derived from caryopses of native Nigerian rice cultivars.⁷⁹ Although *O. glaberrima* and *O. sativa* are both species that are grown by farmers in Nigeria and other African nations, there is much less information available on callus induction and plantlet regeneration of *O. glaberrima*.

A lack of effective in vitro regeneration systems for local cultivars may be the reason why African rice varieties have not been much better utilized with various genetic modifications.^{45,46} For instance, the callus of cultivar IRGC104165 (*O. glaberrima*)⁸⁴ has recently demonstrated a failure of shoot regeneration. The ability to induce and regenerate calluses is also known to be variety-dependent, as it was demonstrated to prevent the genetic modification of many indica rice varieties.⁸⁵ One of the main obstacles to the genetic improvement of this species may be *O. glaberrima*'s poor regeneration capacity caused by a lack of suitable media. However, since a better transformation protocol for *O. glaberrima* landraces can be developed, this shouldn't be a barrier to creating good and acceptable varieties. Testing different hormonal concentration gradients (Kinetin, NAA, or others)⁵⁵ can be used to assess regeneration capacity. Additionally, it has been noted that supplementing media with a substance like a coconut water can enhance the transformation and regeneration of both *O. glaberrima* and *O. sativa* landraces.^{47,55} Moreover, it has been demonstrated that *O. glaberrima* has a greater capacity for regeneration when the concentration of sucrose in the regeneration media is increased.^{63,66}

4. Factors for Improving Plant Regeneration in *O. Glaberrima*

Biotechnologists should first focus more on developing an appropriate transformation and

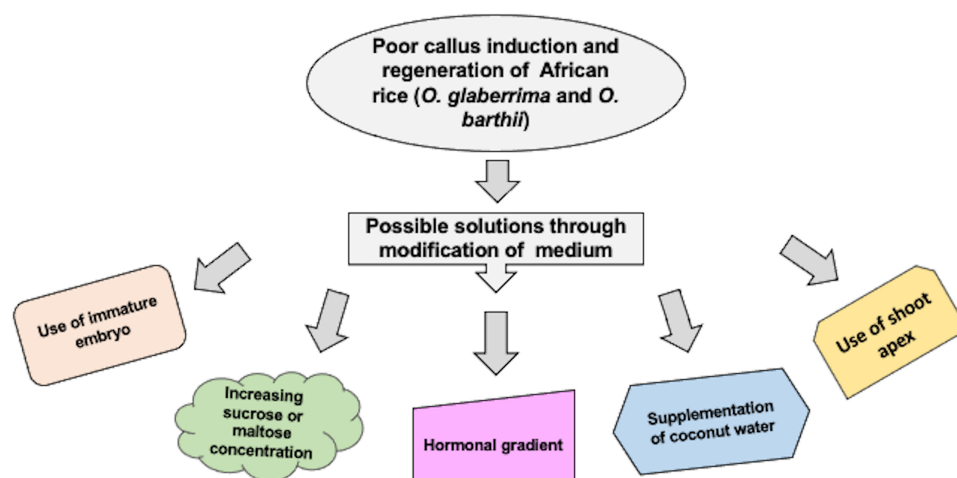


Figure 1. Possible solutions for callus induction and regeneration challenges of African rice *Oryza glaberrima*.

regeneration protocol for *O. glaberrima* to successfully manipulate genetic material and increase rice yield production in Africa. Rather than concentrating on imparting those traits to *O. sativa*, they should focus on improving *O. glaberrima*, which is already adapted to the challenging environmental conditions of Africa. How can we enhance *O. glaberrima*'s transformation and regeneration capacity? The following strategies may be used to enhance African rice's capacity for tissue culture (Figure. 1).

4.1 Modifying Sucrose or Maltose Concentration of Media

Both embryonic development and somatic embryogenic induction are significantly influenced by the type of carbon source and its various concentrations.^{86,87} Since sucrose produced more biomass than fructose and glucose combined, it is known to be the most effective carbon source for causing callus.⁸⁸ However, sucrose has a different rate of organogenesis than maltose and may have a higher osmotic potential, which causes callus browning⁸⁹ and, at concentrations of 6% or higher causing necrosis.⁹⁰ Brisibe et al. study showed that the high frequency of shoot induction in both liquid and agar culture occurred at 5% across the range of sucrose concentrations tested (0–15%).⁶³

Indeed, by increasing the sucrose concentration of MS media from 3 to 5%,⁶⁶ a recent study by Mmbando et al. was able to regenerate transgenic plants from one African rice cultivar, TOG12380

(*O. glaberrima*). However, only one cultivar of the five *O. glaberrima* tested in that study was regenerable. Maltose, on the other hand, has demonstrated superiority for green shoot regeneration in barley⁹¹ when compared to sucrose as a carbon source for androgenesis. According to Sah et al.,⁹² 4% maltose resulted in the highest levels of callus induction (90.33%) and regeneration (82.66%) of the Japanese rice variety Kitaake. In addition, the number of days to callus formation in the media was reduced by maltose (4%) rather than sucrose.⁹² To create the best tissue culture media for this species, we may need to modify and test different maltose concentrations in *O. glaberrima*. Although Mmbando et al.⁶⁶ were able to regenerate *O. glaberrima* at 5% sucrose, the small number of transgenic lines found in this study may be a result of the high sucrose concentration's impact on the regeneration media. Agar concentration changes can also affect the induction and regeneration of calluses.⁹³ Future research should therefore examine the extent of *O. glaberrima* callus induction and regeneration at 4% maltose as well as by varying the type and concentration of agar.

4.2 Hormonal Gradient for Determining Optimum Condition for *O. Glaberrima*

The induction and growth of calluses are known to be significantly influenced by auxin.⁹⁴ Additionally, it has been demonstrated that combining different auxin types is a superior alternative to using a single auxin for the callus induction of rice.⁹⁵ For

instance, a hormonal gradient was created based on the optimal concentration of most *O. glaberrima* accessions, which has been shown to be 0.5–4 mg L⁻¹ kinetin and 0.01–0.05 mg L⁻¹ α -naphthaleneacetic acid (NAA).⁵⁵ The best hormonal setting for NERICA⁹⁶ regeneration has been demonstrated to be 1.3 μ M of NAA and 50 μ M of kinetin. Other studies suggested only using 11.16 μ M of benzylaminopurine (BAA) for *O. glaberrima* regeneration.⁴⁶ Therefore, the success of tissue culture and, consequently, genetic modification of *O. glaberrima* depends on the combination of these growth hormones. However, *O. sativa* is the basis for the majority of the established ideal hormonal conditions, and *O. glaberrima* has no suitable media.⁴⁷ The best conditions for *O. glaberrima* callus induction and regeneration can be discovered by performing a hormonal gradient. Through the use of hormone gradient plates and the addition of coconut water to the growth media, Lacchini et al.⁵⁵ revealed a wide variation in the ability of 13 different African rice accessions to regenerate. In addition, based on earlier studies,⁶⁶ Mmbando et al. tested various hormonal concentrations. They discovered that cultivar TOG12380 (*O. glaberrima*) could regenerate when Brisibe et al. (1990)⁶³ media were given a sucrose concentration increase to 50 g L⁻¹. The MS medium with 10 μ M 2,4-D was deemed suitable for *O. glaberrima* callus induction despite the lack of a hormonal gradient in that study, while the sucrose concentration was increased to 50 g L⁻¹ with 0.1 μ M of 2,4-D and 50 μ M kinetin for regeneration. In the future, it will need to be determined whether using maltose rather than sucrose with the same hormonal combination will produce better results. Additionally, adding a hormonal gradient to the medium, as suggested by Mmbando et al.,⁶⁶ may improve *O. glaberrima*'s ability to grow in tissue culture.

4.3 Supplementation of Coconut Water in Callus Induction Media

A routine tissue culture system that includes callus induction and plantlet regeneration is a basic requirement for the genetic modification and generation of transgenic rice plants. Supplementing with coconut water is one way to ensure that your

calluses are healthy (CW). Rice,^{47,97} and even other species⁹⁸ have demonstrated that CW, the colorless liquid endosperm of green coconuts (*Cocos nucifera*),⁹⁹ increases the size of the calli and regeneration. For example, Mamun et al. (2004) showed that 2, 4-D, and 10% coconut milk produced the highest amount of regenerative callus from leaf sheath in sugarcane.¹⁰⁰ Because coconut water contains more sugars¹⁰¹ and is rich in essential amino acids like methionine, histidine, lysine, and cysteine,¹⁰² it has been shown to improve the ability of tissue culture. Additionally, it contains a lot of minerals, such as potassium, vitamins, magnesium, and calcium,¹⁰³ all of which promote rapid callus initiation. Although coconut milk increased callus size in the majority of cultivars, at least some *O. glaberrima* variety has been reported to not require coconut milk for callus formation.⁴⁷ On the other hand, a recent study by Lacchini et al. used coconut water to enhance the medium and regeneration of African landraces.⁵⁵ Since the study was primarily conducted on African rice landraces, it provided an opportunity to design the best media for African rice. The inclusion of CW should be viewed as a crucial element in creating the best media for African rice, *O. glaberrima*.

4.4 Use of Immature Embryo Instead of Mature Seeds

Immature embryos may be one way to improve *O. glaberrima*'s capacity for transformation because conventional methods for transforming wild *Oryza* accessions typically fail to produce calli from the scutellum tissue of embryos in mature seeds.¹⁰⁴ As demonstrated on *indica* varieties that are resistant to transformation and tissue culture,^{105,106} immature embryos have shown to be a successful method for introducing genes without inducing callus.^{80,105} As recently demonstrated on cultivar IRGC104165 (*O. glaberrima*),⁸⁴ most *O. glaberrima* have poor callus induction, which fails shoot regeneration. Thus, introducing the gene of interest by immature embryos rather than mature seeds might increase the regeneration ability of this species. Although Mmbando et al. attempt to transform the *O. barthii* cultivar (TOB7307) using an immature embryo failed, a study by Shimizu et al.¹⁰⁴ showed that wild *Oryza* could be transformed using

modified immature embryo techniques. The different samples and transformation and regeneration conditions used in these two studies may be to blame for the dissimilar findings. Additionally, aside from the impact of the *Agrobacterium* strains used,⁸⁰ heat treatment and centrifugation before *Agrobacterium* infection can increase transformation efficiency in cultivated maize, wheat, sorghum, and rice.^{105,107} Thus, including these procedures in the *O. glaberrima* embryo at an early stage could improve transformation effectiveness, enable genome editing, and hasten research on other wild *Oryza* genetic resources. In a recent study, Liang et al. improved *Agrobacterium*-mediated transformation and regeneration methods for *indica* rice from immature embryos as explants by disrupting SUB1A as a test case via CRISPR-Cas9.¹⁰⁸ Since *O. glaberrima* and *indica* are both known to be resistant to tissue culture, it may be worthwhile to modify the medium for Ciherang-Sub1 and test it with *O. glaberrima*. Although there are a few exceptions,^{55,66,109} *O. glaberrima* varieties^{45,46,63,84} frequently fail to transform using the mature embryo method; therefore, the immature embryo method may be used on this species. We must not ignore the fact that, despite being a good alternative, this protocol is expensive and necessitates more time and labor to dissect immature embryos.

4. 5 Use of Rice Shoot Apex (Meristem) in Transformation and Regeneration

The ability to induce and regenerate calluses is genotype-dependent, which sets limits on the genetic engineering of *indica* and African rice varieties. The main obstacles to using these explants in transformation include multiple stages of subculture to select the transformed calli, loss of regeneration potential in calli, availability of immature embryos only during a specific timeframe, and difficulties with isolation and sterilization of immature embryos.⁸¹ The main benefit of shoot meristem-based multiplication systems, however, is that they allow for the quick and direct production of transgenic plants from transformed shoot apices⁷⁰ and are typically genotype-independent, making them applicable to various genotypes.⁸¹ As a result, using the rice shoot apex for transformation and recovering *O. glaberrima* through *Agrobacterium*-

mediated transformation could be a potential solution. This approach has proven to be simple to use and capable of preserving cultivar integrity.^{110,111} Changing the growth regulators and carbon source of plants while using the shoot apex may make it possible to create an effective and speedy protocol for *Agrobacterium*-mediated genetic transformation of African rice varieties. The generation of chimeric during regeneration, however, is one of the complications linked to this technique that may impede the development of transgenic plants.¹¹²

5. Enhancing Yield Potential of *O. Glaberrima* through Gene Editing: Target Traits

How to increase food production is a major issue for the development of many African nations. The best way to increase rice production in Africa may therefore be to use modern biotechnology to increase the native African rice variety *O. glaberrima*'s resistance to both biotic and abiotic stresses. Despite negative press in Africa, it has been proposed that biotechnology is a key driver for commercializing crops that provide all with a good standard of living and jobs.¹¹³ Although the issues and difficulties related to biotechnology in Africa will not be the focus of this section, these topics have already been covered elsewhere.^{10,75,76,114–121} Recently, several gene editing methods based on the transcription-activator¹²² like effector nucleases (TALEN), zinc finger nucleases (ZFNs), CRISPR/Cas9 system, and meganucleases have been developed. Agroinfiltration (true agro-infiltration, agro-inoculation or infection, and floral dip), cisgenesis, and grafting (non-genetically modified (GM) scion on GM rootstock or the opposite) are additional new plant breeding techniques (NPBT). The design of a guide RNA (gRNA) for CRISPR genome editing has proven to be simpler than that for TALENs and ZFNs.⁵⁴ The safety of GM crops like rice for human consumption is a major concern, particularly the health risks that could arise from consuming GM foods that contain toxins and allergens.¹²³ With the complexity of the biosafety regulation being reduced,^{54,124} recent gene editing techniques utilizing the CRISPR/Cas 9 protocol have demonstrated an improvement in simplicity, amicability, precision, adaptability, and efficiency in the process of

genetically modified crops. Additionally, it is simpler to implement and calls for creating a guide gRNA of roughly 20 nucleotides that is complementary to the DNA stretch within the target gene that results in the production of accurate insertion or deletion events.^{54,75}

Because the gene to be modified is already present in the plant's DNA,¹²⁵ CRISPR/Cas 9 edited crops have additional advantages over transgenic plants as opposed to the transgenic method, which produced random insertions and phenotypes. Particularly in Africa, where there are many opponents of using GM crops and biotechnology, CRISPR/Cas9 mediated genome editing (CMGE) may present a great opportunity for editing the undesirable traits of *O. glaberrima* with a significant reduction in problems resulting from GM crops. Simple procedures in CMGE make it possible for even a small laboratory with a basic plant transformation setup typically for many lower-income countries in Africa and other developing countries to carry out genome editing projects.⁵⁴ To carry out the CRISPR-Cas9 project, the following steps must be taken: first, locate the protospacer adjacent motif (PAM) sequence in the target gene; second, create a single gRNA (sgRNA); third, clone the sgRNA into an appropriate binary vector; fourth, introduce the sgRNA into host species/cell lines by transformation; fifth, screen; and finally, validate the edited lines.⁵⁴ Although the CMGE method has made it possible to create non-genetically modified (Non-GMO) crops with the desired trait and increased yield under a variety of abiotic and biotic stress conditions, it has been much less utilized to increase *O. glaberrima*'s yield production. The creation of significant off-target cleavage sites has been one of some CRISPR/Cas9 systems' drawbacks. However, among the suggestions for minimizing this issue are the lengthening of the protospacer adjacent motif⁵⁴ and the use of a more precise gRNAs design approach.¹²⁶ But the question is, can Africa benefit from these new gene editing techniques? Can these techniques enhance *O. glaberrima*'s nutritional value and yield, which is underutilized but important for both local African farmers and the rest of the world? In this study, several methods are suggested for removing all undesirable traits from *O. glaberrima* by CMGE.

This study suggested that the CMGE technique could be used to increase the yield of *O. glaberrima* and even accelerate its domestication,^{55,127–132} similar to the increase in yield in *O. sativa* caused by the simultaneous introduction of three trait-related quantitative trait loci (QTLs)¹³³ Since the publication of the *O. glaberrima* genome in 2014,⁷² genes for numerous agronomic traits in rice, including *GRAIN WEIGHT 2* (GW2), *GRAIN NUMBER 1A* (GN1A), and *GRAIN SIZE 3* (GS3), have already been reported.¹³⁴ By stacking numerous unfavorable traits among this species, it opens the door for yield improvements in *O. glaberrima*.^{55,135} *O. glaberrima*'s main drawbacks¹³⁶ include lodging, low yield, grain shattering,⁸ and possibly UVB sensitivity.¹⁷ The inexpensive and straightforward method of CRISPR/Cas9 may provide a great opportunity to increase the yield of this species by carrying out multiple knockouts of the genes encoding for these traits in *O. glaberrima*. Instead of being the sole genetic resource for improving *O. sativa*,¹⁸ which cannot survive in Africa's harsh environment and results in low production for local farmers who depend on rice as a source of income, this will quicken its breeding programs. A further method for increasing the yield of *O. glaberrima* might involve editing the yield genes grain number 1a (*Gn1a*) and *DENSE AND ERECT PANICLE1* (DEP1), which are important contributors to rice yield increases. Genes like *OsGRAS19* and *D26*,¹³⁷ which control *O. sativa* grain size as well, have the added benefit of enhancing the grain quality of this species. Additionally, the high UVB sensitivity of native African landraces could be reduced by inhibiting the downstream molecular pathways, such as growth-regulatory factors, microRNA396 or hormone-associated genes^{138,139} or other transcription factors, that cause UVB-induced leaf growth inhibition of cell proliferation.

Since the genes that act as a barrier to interspecific hybridization for breeding programs of the two species have already been identified,^{50,51,140} it is possible to use the CRISPR/Cas9 technique to eliminate these genes in *O. glaberrima*. We cannot, however, rule out the possibility that *O. glaberrima* would not survive in the harsh environment of Africa if all of these genes were removed. Many

essential genes must be overexpressed for many agriculturally significant traits, such as increased photosynthesis, to occur, as their knockout causes seedling lethality.¹⁴¹ Therefore, before approving modified plants, field trials evaluating the efficacy and lethality of these multiple knockout-generated mutants should be carefully assessed. The CRISPR/Cas9 technique has also been used to induce resistance to bacterial leaf blight (BLB), one of the most harmful vascular diseases of rice; which is caused by the proteobacterium *Xanthomonas oryzae* pv. *oryzae* (*Xoo*) and causes 75% of yield loss with millions of hectares annually.¹⁴² The sugar-transporting SWEET gene family in rice is typically the target of transcription activator-like effectors (TALEs), which are primarily in charge of releasing the sugar into the apoplast for the nutrition of the *Xoo* pathogens.¹⁴³ For instance, using the CRISPR/Cas9 technique, the introduction of InDels into the effector-binding element (EBE) of the OsSWEET13 promoter of the MS14K line and the creation of a novel rice line with three edited OsSWEET EBEs led to broad-spectrum resistance against all of the tested *Xoo* strains.¹⁴⁴ Additionally, Olivia et al.¹⁴⁵ introduced mutations at the promoters of all three SWEET genes at EBEs recognized by different *Xoo* TALEs using CRISPR/Cas9. This resulted in strong and widespread BLB resistance. A CRISPR/Cas9 targeted mutation in the ethylene-responsive factor, OsERF922 in rice,¹⁴⁶ significantly increased the resistance to blast disease caused by *Magnaporthe oryzae*.

Similar results have been seen when the MLO allele has been eliminated in tomato and wheat.^{147,148} Even though the majority of *O. glaberrima* varieties have demonstrated resistance to certain diseases, such as the rice yellow mottle virus,^{36,37} the CMGE technique may still be able to help the susceptible lines increase some biotic resistance, as it has been done with other crops.^{149,150} Furthermore, *O. glaberrima* genes have been used to enhance particular grain quality traits in high-yielding *O. sativa* hybrid cultivars.⁴³ The development of strategies for the increased milling and cooking of African rice has been made easier by QTL mapping grain quality traits, particularly *hr3*, *Amy6*, and *Alk6-1* from an interspecific cross between *Oryza sativa* and *Oryza glaberrima*.⁴²

Through non-host gene insertion in *O. sativa*, transgenic rice with a variety of agronomically significant traits, such as seed storage protein Bean - phaseolin,¹⁵¹ pea legumin,¹⁵² soybean glycinin,¹⁵³ and daffodil phytoene synthase, provitamin A,¹⁵⁴ has already been produced. These traits offer opportunities to improve the nutritional^{155,156} quality of *O. glaberrima* by CMGE. Because *O. sativa* and *O. glaberrima* are both growing in SSA, increasing the production of both species using CRISPR/Cas9 techniques as well as genetic engineering is crucial for feeding the world's expanding population, especially Africa, which is the hotspot. Since rice is used as both a food and a cash crop, increasing rice production will also help poor farmers' economies.

6. Conclusion

African rice varieties^{45,46,63,84} and *indica*^{105,106} are both known to be resistant to tissue culture and transformation, which impedes genetic modification methods. Therefore, it is essential to develop a suitable transformation and regeneration protocol to unlock all of these species' advantageous traits, which could significantly improve food security. The highly complex process of biotechnology necessitates a certain critical mass of technical, financial, and intellectual resources. African governments must educate their people about the advantages of biotechnology for agriculture, industry, and health care, and develop strong policies to support emerging gene-editing technologies. The development of global regulatory frameworks for gene-edited crops is necessary before the implementation of CRISPR technology for food crops, particularly in Africa, where there are many anti-biotechnologists.^{75,76,115,118,119} Due to the importance of these molecular biology techniques, African governments must give science a higher priority in their budgets. To do this, they must train more biotechnologists and set up contemporary labs with sufficient molecular biology reagents and equipment. The pyramiding of genes for the improvement of *O. glaberrima* can also make use of the abundance of rice sequencing data and the accessibility of numerous QTL markers. On the other hand, African researchers and scientists need to take an interest in enhancing

O. glaberrima and form partnerships with universities and biotech companies in developed countries. It is now time for Africa to develop the skills necessary to feed her population using her resources, including professionals and crops like rice (*O. glaberrima*). The mind-set of the majority of African scientists must be changed in order for this to be accomplished. *O. glaberrima* is still grown in many regions of SSA by local farmers.⁹ Local farmers still prefer *O. glaberrima* over many improved *O. sativa* and NERICA varieties,¹⁰ which shows that breeders need to pay more attention to this underappreciated but crucial species for SSA. Local farmers in SSA have the chance to increase the yield and nutritional value of *O. glaberrima* by using contemporary genetic editing tools like CRISPR-Cas9. Therefore, this study put forth the hypothesis that genome editing techniques may be an ideal tool for eradicating those undesirable traits through the insertion of non-host genes, regulation of transcription or translation, and modification of host sequences, rather than ignoring this species with economically significant traits for poor farmers in SSA. Future research should investigate whether altering these genes in *O. glaberrima* could affect the organism's resistance to additional stresses on the continent of Africa.

This study highlighted the advantages and disadvantages of *O. glaberrima* relative to *O. sativa* and the necessity of enhancing this species for high yield production in SSA. This study also offers some recommendations for improving *O. glaberrima*'s capacity for regeneration. To fully utilize the benefits of molecular breeding techniques on this species, this study urged the urgent development of an effective transformation and regeneration protocol for *O. glaberrima* and other African landraces. This study offers suggestions for enhancing *O. glaberrima* breeding and yield to feed the expanding global population, particularly in Africa.

Possible ways for having high transformation and regeneration in African rice varieties

Acknowledgments

The author enthusiastically acknowledges UDOM for providing office space for research write-ups.

Author Contribution Statement

G.S.M: Conceived, designed, interpreted the data, and wrote the paper.

Disclosure Statement

No potential conflict of interest was reported by the author(s).

Funding

This research did not receive any specific grant from funding agencies in the public, commercial, or not-for-profit sectors.

ORCID

Gideon Sadikiel Mmbando  <http://orcid.org/0000-0001-8014-9072>

References

1. Hamilton RW, Ammayao A. The art of rice: spirit and sustenance in Asia. US: University of California Los Angeles, Fowler Museum of Cultural History; 2003.
2. FAO. 2012. State of food and agriculture 2012: investing in agriculture for a better future. FAO.
3. Jung K-H, An G, Ronald PC. Towards a better bowl of rice: assigning function to tens of thousands of rice genes. *Nat Rev Genet*. 2008;9:91–101.
4. Vaughan DA, Morishima H, Kadowaki K. Diversity in the *Oryza* genus. *Curr Opin Plant Biol*. 2003;6:139–46.
5. Jones MP, Dingkuhn M, Aluko GK, Semon M. Interspecific *Oryza sativa* L. X *O. glaberrima* Steud. progenies in upland rice improvement. *Euphytica*. 1997;94:237–46.
6. Londo JP, Chiang Y-C, Hung K-H, Chiang T-Y, Schaal BA. Phylogeography of Asian wild rice, *Oryza rufipogon*, reveals multiple independent domestications of cultivated rice, *Oryza sativa*. *Proc Natl Acad Sci*. 2006;103:9578–83.
7. Van Andel T. African rice (*Oryza glaberrima* Steud.): lost crop of the enslaved Africans discovered in Suriname. *Econ Bot*. 2010;64:1–10.
8. Linares OF. African rice (*Oryza glaberrima*): history and future potential. *Proc Natl Acad Sci*. 2002;99:16360–65.
9. Singh BN. Rice Growing environments and biophysical constraints in different ecological zones in Nigeria Met. J. 1997;2:35–44.
10. Sié M, Sanni K, Futakuchi K, Manneh B, Mandé S, Vodouhe RS, Dogbe S, Drame K, Ogunbayo A, Ndjiondjop M-N. Towards a rational use of African rice (*Oryza glaberrima* Steud.) for breeding in Sub-Saharan Africa Genes, gen. genom. 2012;5

- (1):81–90. doi:[10.1161/CIRCGENETICS.111.959817](https://doi.org/10.1161/CIRCGENETICS.111.959817).
11. Futakuchi K, Fofana M, Sié M. Varietal differences in lodging resistance of African rice (*Oryza glaberrima* Steud.). *Asian J Plant Sci*. 2008;7(6):569–73. doi:[10.3923/ajps.2008.569.573](https://doi.org/10.3923/ajps.2008.569.573).
 12. Sié M, Dogbe SY, Coulibaly M. Selection of interspecific hybrids (*O. sativa* x *O. glaberrima* or lowland NERICAs) and intraspecifics adapted to rainfed lowland growing conditions. *Int Rice Comm Newslett*. 2005;54:47–51.
 13. Somado EA, Guei RG, Keya SO. . 2008 NERICA: the new rice for Africa: a compendium (Cote, d'Ivoire, Abidjan) pp. 10–14.
 14. Futakuchi K, Sié M. Better exploitation of African rice (*Oryza glaberrima* Steud.) in varietal development for resource-poor farmers in West and Central Africa. *Agric J*. 2009;4:96–102.
 15. Koffi G. Collection and conservation of existing rice species and varieties of Africa. *Agron Trop*. 1980;34:228–37.
 16. Dingkuhn M, Jones MP, Johnson DE, Sow A. Growth and yield potential of *Oryza sativa* and *O. glaberrima* upland rice cultivars and their interspecific progenies. *F Crop Res*. 1998;57:57–69.
 17. Mmbando GS, Teranishi M, Hidema J. Very high sensitivity of African rice to artificial ultraviolet-B radiation caused by genotype and quantity of cyclobutane pyrimidine dimer photolyase. *Sci Rep*. 2020;10:1–14.
 18. Sarla N, Swamy BPM. *Oryza glaberrima*: a source for the improvement of *Oryza sativa*. *Curr Sci*. 2005;89:955–63.
 19. Sakagami J-I, Kawano N. Survival of submerged rice in a flood-prone region of West Africa. *Tropics*. 2011;20:55–66.
 20. Jones PM, Mande S, Aluko K. Diversity and potential of *Oryza glaberrima* Steud in upland rice breeding. *Japanese J Breed*. 1997;47:395–98.
 21. Tobita S, Sahrawat KL, Diatta S, Jones MP Response of African rice (*Oryza glaberrima* Steud.) to phosphate application in the upland of a P-deficient soil in the humid forest zone of West Africa. In: *Proceedings of the 2nd International Symposium on Phosphorus Dynamics in the Soil-Plant Continuum*, Perth, Western Australia. 2003; 21–26.
 22. Futakuchi K, Jones MP, Ishii R. Physiological and morphological mechanisms of submergence resistance in African rice (*Oryza glaberrima* Steud.). *Japanese J Trop Agric*. 2001;45:8–14.
 23. Kawano N, Ito O, Sakagami J-I. Flash flooding resistance of rice genotypes of *Oryza sativa*. *Environ Exp Bot*. 2008;63:9–18.
 24. Maji AT, Gana AS, Ukwungwu MN. Responses of *Oryza glaberrima* accessions to rice stresses and their morphological characteristics. *African J Gen Agric*. 2011;6:732–37.
 25. Ndjiondjop MN, Cisse F, Girma G, Sow M, Djedatin G, Blandine F. Morpho-agronomic and molecular characterisation of *Oryza glaberrima* germplasm from Mali. *J Biotechnol*. 2010;9:7409–17.
 26. Sahrawat KL, Sika M. Comparative tolerance of *Oryza sativa* and *O. glaberrima* rice cultivars for iron toxicity in West Africa. *Int Rice Res Notes*. 2002;27:30–31.
 27. Coyne DL, Plowright RA, Fofana B. Observations on the susceptibility of *Oryza sativa* and resistance of *Oryza glaberrima* to the cyst nematode (*Heterodera sacchari*) and the influence of weed management in upland rice in Ivory Coast. *Int J Pest Manag*. 1999;45:255–58.
 28. Plowright RA, Coyne DL, Nash P, Jones MP. Resistance to the rice nematodes *Heterodera sacchari*, *Meloidogyne graminicola* and *M. incognita* in *Oryza glaberrima* and *O. glaberrima* x *O. sativa* interspecific hybrids. *Nematology*. 1999;1:745–51.
 29. Soriano I, Schmit V, Brar D, Prot J-C, Reversat G. Resistance to rice root-knot nematode *Meloidogyne graminicola* identified in *Oryza longistaminata* and *O. glaberrima*. *Nematology*. 1999;1:395–98.
 30. Lorieux M, Reversat G, Diaz SXG, Denance C, Jouvenet N, Orioux Y, Bourger N, Pando-Bahuon A, Ghesquière A. Linkage mapping of Hsa-1 Og, a resistance gene of African rice to the cyst nematode, *Heterodera sacchari*. *Theor Appl Genet*. 2003;107:691–96.
 31. Ukwungwe MN, Williams CT, Okhidlevble O. Screening of African rice, *Oryza glaberrima* Steud., for resistance to the African rice gall midge *Orseolia oryzivora* Harris and Gagne. *J Food Technol Africa*. 1999;4:3.
 32. Williams CT. Multilocal screening of *Oryza sativa* and *O. glaberrima* for resistance to African rice gall midge *Orseolia oryzivora* in West Africa. *Int Rice Res Notes*. 1999;24:26–27.
 33. Johnson DE, Dingkuhn M, Jones MP, Mahamane MC. The influence of rice plant type on the effect of weed competition on *Oryza sativa* and *Oryza glaberrima*. *Weed Res*. 1998;38:207–16.
 34. Dingkuhn M, Johnson DE, Sow A, Audebert AY. Relationships between upland rice canopy characteristics and weed competitiveness. *F Crop Res*. 1999;61:79–95.
 35. Fofana B, Rauber R. Weed suppression ability of upland rice under low-input conditions in West Africa. *Weed Res Oxford*. 2000;40:271–80.
 36. Abo ME, Sy AA, Alegbejo MD. Rice yellow mottle virus (RYMV) in Africa: evolution, distribution, economic significance on sustainable rice production and management strategies. *J Sustain Agric*. 1998;11:85–111.
 37. Ndjiondjop M-N, Albar L, Fargette D, Fauquet C, Ghesquière A. The genetic basis of high resistance to rice yellow mottle virus (RYMV) in cultivars of two cultivated rice species. *Plant Dis*. 1999;83:931–35.
 38. Mohapatra S. Pockets of gold. *Rice Today*. 2010;9:32–33.
 39. Teeken B African rice (*Oryza glaberrima*) cultivation in the Togo Hills: ecological and socio-cultural cues in

- farmer seed selection and development. PhD diss., Wageningen University and Research, 2015.
40. Ricachenevsky FK, Sperotto RA. Into the wild: *Oryza* species as sources for enhanced nutrient accumulation and metal tolerance in rice. *Front Plant Sci.* 2016;7:974.
 41. Martínez CP, Borrero J, Carabali SJ, Delgado D, Correa F, Tohme J High-iron and zinc rice lines for Latin America. In: Proc. 31st Rice Tech. High-iron and zinc rice lines for Latin America [poster][on line], Cali, Colombia. 2006.
 42. Aluko G, Martinez C, Tohme J, Castano C, Bergman C, Oard JH. QTL mapping of grain quality traits from the interspecific cross *Oryza sativa* x *O. glaberrima*. *Theor Appl Genet.* 2004;109:630–39.
 43. Li J, Xiao J, Grandillo S, Jiang L, Wan Y, Deng Q, Yuan L, McCouch SR. QTL detection for rice grain quality traits using an interspecific backcross population derived from cultivated Asian (*O. sativa* L.) and African (*O. glaberrima* S.) rice. *Genome.* 2004;47:697–704.
 44. Watanabe H. Grain protein content of African rice (*Oryza glaberrima* Steud.) lines and Asian rice (*O. sativa* L.) varieties with West Africa. *Oryza.* 2004;41:35–38.
 45. Brisibe EA, Miyake H, Taniguchi T, Maeda E. Callus formation and scanning electron microscopy of plantlet regeneration in African rice (*Oryza glaberrima* Steud). *Plant Sci.* 1992;83:217–24.
 46. Diawuoh RG, Klu GYP, Amoatey HM, Kusi-adjei R, Quartey EK. Callus Induction and plant regeneration from dehusked mature seeds of three accessions of African rice (*Oryza glaberrima* Steud). *J Biol Agric Heal.* 2016;6:31–39.
 47. Fatokun CA, Yamada Y. Variations in callus formation and plant regeneration in African rice (*Oryza glaberrima* Steud). *J Plant Physiol.* 1984;117:179–83.
 48. Sano Y. 1986. Sterility barriers between *Oryza sativa* and *O. glaberrima* In Rice Genetics 1 (In 2 parts) Rice Genetics Proceedings of the First Rice Genetics Symposium (Manila: IRR1) p. 109–18.
 49. Kato S. On the affinity of rice varieties as shown by fertility of hybrid plants. *Bull Sci Fac Agric Kyushu Univ.* 1928;3:132–47.
 50. Koide Y, Ogino A, Yoshikawa T, Kitashima Y, Saito N, Kanaoka Y, Onishi K, Yoshitake Y, Tsukiyama T, Saito H, et al. Lineage-specific gene acquisition or loss is involved in interspecific hybrid sterility in rice. *Proc Natl Acad Sci U S A.* 2018;115:E1995–E1962.
 51. Xie Y, Xu P, Huang J, Ma S, Xie X, Tao D, Chen L, Liu YG. Interspecific hybrid sterility in rice Is mediated by OgTPR1 at the S1 locus encoding a peptidase-like protein. *Mol Plant.* 2017;10:1137–40.
 52. Bortesi L, Fischer R. The CRISPR/Cas9 system for plant genome editing and beyond. *Biotechnol Adv.* 2015;33:41–52.
 53. Hsu PD, Scott DA, Weinstein JA, Ran FA, Konermann S, Agarwala V, Li Y, Fine EJ, Wu X, Shalem O. DNA targeting specificity of RNA-guided Cas9 nucleases. *Nat Biotechnol.* 2013;31:827–32.
 54. Jaganathan D, Ramasamy K, Sellamuthu G, Jayabalan S, Venkataraman G. CRISPR for crop improvement: an update review. *Front Plant Sci.* 2018;9:1–17.
 55. Lacchini E, Kiegle E, Castellani M, Adam H, Jouannic S, Gregis V, Kater MM. CRISPR-mediated accelerated domestication of African rice landraces. *PLoS One.* 2020;15:e0229782.
 56. Futakuchi K, Sié M, Saito K. Yield potential and physiological and morphological characteristics related to yield performance in *Oryza glaberrima* Steud. *Plant Prod Sci.* 2012;15:151–63.
 57. Komari T, Hiei Y, Ishida Y, Kumashiro T, Kubo T. Advances in cereal gene transfer. *Curr Opin Plant Biol.* 1998;1:161–65.
 58. Tanksley SD, McCouch SR. Seed banks and molecular maps: unlocking genetic potential from the wild. *Science.* 1997;277:1063–66.
 59. Datta SK. Rice biotechnology: a need for developing countries. *AgBioForum.* 2004;7:31–35.
 60. Dievart A, Perin C, Hirsch J, Bettembourg M, Lanau N, Artus F, Bureau C, Noel N, Droc G, Peyramard M. The phenome analysis of mutant alleles in leucine-rich repeat receptor-like kinase genes in rice reveals new potential targets for stress tolerant cereals. *Plant Sci.* 2016;242:240–49.
 61. Murashige T, Skoog F. A revised medium for rapid growth and bio assays with tobacco tissue cultures. *Physiol Plant.* 1962;15:473–97.
 62. Lee S-T, Huang W-L. Cytokinin, auxin, and abscisic acid affects sucrose metabolism conduce to de novo shoot organogenesis in rice (*Oryza sativa* L.) callus. *Bot Stud.* 2013;54:1–11.
 63. Brisibe EA, Taniguchi T, Maeda E. In-vitro plant regeneration from morphogenic callus cultures of cultigens and wild *Oryza* species. *Japanese J Crop Sci.* 1990;59:557–65.
 64. Chu -C-C, Wang -C-C, Sun C-S, C HSU, K-c YIN, F-y BI. Establishment of an efficient medium for anther culture of rice through comparative experiments on the nitrogen sources. *Sci Sin.* 1975;18:659–68.
 65. Al-Forkan M, Rahim MA, Chowdhury T, Akter P, Khalela L. Development of highly *in vitro* callogenesis and regeneration system for some salt tolerant rice (*Oryza sativa* L.) cultivars of Bangladesh. *Biotechnology.* 2005;4:230–34.
 66. Mmbando GS, Teranishi M, Hidema J. Transgenic rice *Oryza glaberrima* with higher CPD photolyase activity alleviates UVB-caused growth inhibition. *GM Crop Food.* 2021;12:435–48.
 67. Meneses A, Flores D, Muñoz M, Arrieta G, Espinoza AM. Effect of 2, 4-D, hydric stress and light on *indica* rice (*Oryza sativa*) somatic embryogenesis. *Rev Biol Trop.* 2005;53:361–68.
 68. He Y, Guo X, Lu R, Niu B, Pasapula V, Hou P, Cai F, Xu Y, Chen F. Changes in morphology and biochemical indices in browning callus derived from *Jatropha curcas* hypocotyls. *Plant Cell, Tissue Organ Cult.* 2009;98:11–17.

69. Zuraida AR, Naziah B, Zamri Z, Sreeramanan S. Efficient plant regeneration of Malaysian *indica* rice MR 219 and 232 via somatic embryogenesis system. *Acta Physiol Plant*. **2011**;33:1913–21.
70. Dey M, Bakshi S, Galiba G, Sahoo L, Panda SK. Development of a genotype independent and transformation amenable regeneration system from shoot apex in rice (*Oryza sativa* spp. *indica*) using TDZ. **2012**;2:233–40.
71. Kennedy D. The importance of rice. *Science*. **2002**;296:13.
72. Wang M, Yu Y, Haberer G, Marri PR, Fan C, Goicoechea JL, Zuccolo A, Song X, Kudrna D, Ammiraju JSS, et al. The genome sequence of African rice (*Oryza glaberrima*) and evidence for independent domestication. *Nat Genet*. **2014**;46:982–88.
73. Bakshi S, Dewan D. Status of transgenic cereal crops: a review. *Clon Transgen*. **2013**;3:2.
74. Keener K, Hoban T, Balasubramanian R. Biotechnology and it applications. North Carolina State Univ Coll Agric Life Sci Work Pap March. **2013**;16.
75. Zenna N, Manneh B, Atnaf M, Worede F. Molecular breeding and biotechnology for rice improvement in the developing world, Mulugeta, F, Dawit, A, Milahum, F, Kiyoshi, S (Addis Ababa, Ethiopi). **2019** Advnce in Rice Research and Development in Ethiopia; 79–98. EIAR.
76. Wambugu F. Why Africa needs agricultural biotech. *Nature*. **1999**;400:15–16.
77. Zenna N, Senthilkumar K, Sie M. Rice production in Africa. In: Rice Production Worldwide, Chauhan, BS, Jarban, K, Mahajan, G. Switzerland: Springer; **2017**. p. 117–35.
78. Yookongkaew N, Srivatanakul M, Narangajavana J. Development of genotype-independent regeneration system for transformation of rice (*Oryza sativa* ssp. *indica*). *J Plant Res*. **2007**;120:237–45.
79. Afolabi AS, Oyebanji O, Odusanya O, Abo ME, Misra M, Ogbadu GH. Regeneration of plants from rice caryopsis derived callus culture of Nigerian local cv Suakoko 8 and a NERICA cv FARO 55. *Int J Plant Breed Genet*. **2019**;2:001–4.
80. Hiei Y, Komari T. *Agrobacterium*-mediated transformation of rice using immature embryos or calli induced from mature seed. *Nat Protoc*. **2008**;3:824–34.
81. Kishore SN, Visarada K, Lakshmi YA, Pashupatinath E, Rao SV, Seetharama N. *In vitro* culture methods in sorghum with shoot tip as the explant material. *Plant Cell Rep*. **2006**;25:174–82.
82. Yu K, Pauls KP. Identification of a RAPD marker associated with somatic embryogenesis in alfalfa. *Plant Mol Biol*. **1993**;22:269–77.
83. Takeuchi Y, Abe T, Sasahara T. RFLP mapping of QTLs influencing shoot regeneration from mature seed-derived calli in rice. *Crop Sci*. **2000**;40:245–47.
84. Hu M, Lv S, Wu W, Fu Y, Liu F, Wang B, Li W, Gu P, Cai H, Sun C, et al. The domestication of plant architecture in African rice. *Plant J*. **2018**;94:661–69.
85. Cheema AA, Rashid M, Ashraf M, Qamar ZU. Genetic divergence in rice collections. *Pakistan J Bot*. **2004**;36:557–66.
86. Gerdakaneh M, Mozafari AA, Khalighi A, Sioseh-Mardah A. The effects of carbohydrate source and concentration on somatic embryogenesis of strawberry (*Fragaria* × *ananassa* Duch.). *Am-Eurasian J Agric Env Sci*. **2009**;6:76–80.
87. Thakur T, Sinha K, Kaur T, Kapoor R, Kumar G, Bhunia RK, Salvi P. Efficient genetic transformation of rice for CRISPR/Cas9 mediated genome-editing and stable overexpression studies: a case study on rice lipase 1 and galactinol synthase encoding genes. *Agronomy*. **2022**;12(1).
88. Jayaraman S, Daud NH, Halis R, Mohamed R. Effects of plant growth regulators, carbon sources and pH values on callus induction in *Aquilaria malaccensis* leaf explants and characteristics of the resultant calli. *J For Res*. **2014**;25:535–40.
89. Lentini Z, Reyes P, Martínez CP, Roca WM. Androgenesis of highly recalcitrant rice genotypes with maltose and silver nitrate. *Plant Sci*. **1995**;110:127–38.
90. Grewal D, Gill R, Gosal SS. Factors enhancing induction of high frequency plant regeneration from somatic embryos of *indica* rice (*Oryza sativa* L.). *J Biol Sci*. **2005**;5:697–702.
91. Kuhlmann U, Foroughi-Wehr B. Production of doubled haploid lines in frequencies sufficient for barley breeding programs. *Plant Cell Rep*. **1989**;8:78–81.
92. Sah SK, Kaur A, Sandhu JS. High frequency embryogenic callus induction and whole plant regeneration in *japonica* rice Cv. Kitaake *J Rice Res*. **2014**;2:1–5.
93. Zaidi MA, Narayanan M, Sardana R, Taga I, Postel S, Johns R, McNulty M, Mottiar Y, Mao J, Loit E. Optimizing tissue culture media for efficient transformation of different *indica* rice genotypes. *Agron Res*. **2006**;4:563–75.
94. Nic-Can GI, Loyola-Vargas VM. The role of the auxins during somatic embryogenesis. In: Somatic embryogenesis: fundamental aspects and applications Loyola-Vargas VM, Ochoa-Alejo N. Cham, Switzerland: Springer; **2016**. p. 171–82. 9783319337050.
95. Ge X, Chu Z, Lin Y, Wang S. A tissue culture system for different germplasms of *indica* rice. *Plant Cell Rep*. **2006**;25(5):392–402. doi:10.1007/s00299-005-0100-7.
96. Ishizaki T, Kumashiro T. Genetic transformation of NERICA, interspecific hybrid rice between *Oryza glaberrima* and *O. sativa*, mediated by *Agrobacterium tumefaciens*. *Plant Cell Rep*. **2008**;27(2):319–27. doi:10.1007/s00299-007-0465-x.
97. Biswas A, Mandal AB. Plant regeneration in different genotypes of *indica* rice Indian J Biotech 6 532–40 . **2007**.
98. Al-Khayri JM, Huang FH, Morelock TE, Busharar TA. Spinach tissue culture improved with coconut water.

- HortScience. 1992;27(4):357–58. doi:10.21273/HORTSCI.27.4.357.
99. Abdul-Qadir LH. Callus induction and somatic embryogenesis of rice (*Oryza sativa* L.) improvement with the addition of coconut water Journal of Biology, Agriculture and Healthcare. 2016;6:2.
 100. Mamun MA, Sikdar MBH, Paul DK, Rahman MM, Islam MRI. *In vitro* micropropagation of some important sugarcane varieties of Bangladesh. Asian Journal of Plant Sciences. 2004;3(6):666–69. doi:10.3923/ajps.2004.666.669.
 101. Jackson JC, Gordon A, Wizzard G, McCook K, Rolle R. Changes in chemical composition of coconut(*Cocos nucifera*) water during maturation of the fruit. J Sci Food Agric. 2004;84(9):1049–52. doi:10.1002/jsfa.1783.
 102. Thio GL. Small scale and home processing of fresh coconut (oil manufacture) and utilization of by-products. Amsterdam (Netherlands): Bulletin-Royal Tropical Institute; 1982.
 103. Gopikrishna V, Baweja PS, Venkateshbabu N, Thomas T, Kandaswamy D. Comparison of coconut water, propolis, HBSS, and milk on PDL cell survival J Endod. 34 5 . 2008;587–89.
 104. Shimizu-Sato S, Tsuda K, Nosaka-Takahashi M, Suzuki T, Ono S, Ta KN, Yoshida Y, Nonomura K-I, Sato Y. *Agrobacterium*-mediated genetic transformation of wild *Oryza* species using immature embryos. Rice. 2020;13(1):13. doi:10.1186/s12284-020-00394-4.
 105. Hiei Y, Komari T. Improved protocols for transformation of *indica* rice mediated by *Agrobacterium tumefaciens*. Plant Cell Tissue Organ Cult. 2006;85(3):271–83. doi:10.1007/s11240-005-9069-8.
 106. Hiei Y, Ishida Y, Kasaoka K, Komari T. Improved frequency of transformation in rice and maize by treatment of immature embryos with centrifugation and heat prior to infection with *Agrobacterium tumefaciens*. Plant Cell Tissue Organ Cult. 2006;87(3):233–43. doi:10.1007/s11240-006-9157-4.
 107. Sato-Izawa K, Tokue K, Ezura H. Development of a stable *Agrobacterium*-mediated transformation protocol for *Sorghum bicolor* Tx430. Plant Biotechnol. 2018;35(2):181–85. doi:10.5511/plantbiotechnology.18.0424a.
 108. Liang Y, Biswas S, Kim B, Bailey-Serres J, Septiningsih EM. Improved transformation and regeneration of *indica* rice: disruption of SUB1A as a test case via CRISPR-Cas9. Int J Mol Sci. 2021;22(13):6989. doi:10.3390/ijms22136989.
 109. Lv S, Wu W, Wang M, Meyer RS, Ndjioudjop M-N, Tan L, Zhou H, Zhang J, Fu Y, Cai H, et al. Genetic control of seed shattering during African rice domestication. Nat Plants. 2018;4(6):1049–52. doi:10.1038/s41477-018-0164-3.
 110. Fook CWK, Lai LK, Wong WM, Mahmood M. Efficient regeneration and *Agrobacterium*-mediated transformation protocol for recalcitrant indica rice (*Oryza sativa* L.). Emirates J Food Agric. 2015;27(11):837–48. doi:10.9755/ejfa.2015-08-693.
 111. Clement WKF, Lai KS, Wong MY, Maziah M. Heat and hydrolytic enzymes treatment improved the *Agrobacterium*-mediated transformation of recalcitrant *indica* rice (*Oryza sativa* L.). Plant Cell, Tissue and Organ Culture (PCTOC). 2016;125(1):183–90. doi:10.1007/s11240-015-0926-9.
 112. Raldugina GN, Hoang TZ, Ngoc HB, Karpichev IV. An increased proportion of transgenic plants in the progeny of rapeseed (*Brassica napus* L.) transformants. Vavilov J Genet Breed. 2021;25(2):147. doi:10.18699/VJ21.018.
 113. Brink JA, Woodward BR, DaSilva EJ. Plant biotechnology: a tool for development in Africa. Electron J Biotechnol. 1998;1(2):61–76. doi:10.2225/vol1-issue3-fulltext-6.
 114. Mtui GYS. Status of biotechnology in Eastern and Central Africa. Biotechnol Mol Biol Rev. 2011;6:183–98.
 115. Bornman CH, Grace OM, Van Staden J. Sustainable biotechnology for sub-Saharan Africa: can it be implemented and maintained? South African J Bot. 2004;70(1):1–11. doi:10.1016/S0254-6299(15)30261-1.
 116. Leonce D, Clement I, Sonia U. Plant biotechnology: a key tool to improve crop production in Rwanda. African J Biotechnol. 2019;18(3):68–76. doi:10.5897/AJB2018.16662.
 117. Machuka J. Agricultural Biotechnology for Africa African scientists and farmers must feed their own people. Plant Physiology. 2001;126(1):357–58. doi:10.1104/pp.126.1.16.
 118. Thomson JA. The role of biotechnology for agricultural sustainability in Africa. Philos Trans R Soc B Biol Sci. 2008;363(1492):905–13. doi:10.1098/rstb.2007.2191.
 119. Schurman R. Building an alliance for biotechnology in Africa. J Agrar Chang. 2017;17(3):441–58. doi:10.1111/joac.12167.
 120. FAO CR, Mccouch SR, Herdt RW FAO Rice Conference, Rome, Italy, 12-13 February 2004.
 121. Oja CP, Bull FC, Fogelholm CM, Martin BW. Physical activity recommendations for health: what should Europe do? BMC Public Health. 2010;10:10. doi:10.1186/1471-2458-10-10.
 122. Fraiture M-A, Roosens NHC, Taverniers I, De Loose M, Deforce D, Herman P. Biotech rice: current developments and future detection challenges in food and feed chain. Trends Food Sci Technol. 2016;52:66–79. doi:10.1016/j.tifs.2016.03.011.
 123. Bajaj S, Mohanty A. Recent advances in rice biotechnology-towards genetically superior transgenic rice. Plant Biotechnol J. 2005;3(3):275–307. doi:10.1111/j.1467-7652.2005.00130.x.
 124. Mishra R, Zheng W, Joshi RK, Kaijun Z. Genome editing strategies towards enhancement of rice disease resistance Rice Sci. 2021;28:133–45.
 125. Malzahn A, Lowder L, Qi Y. Plant genome editing with TALEN and CRISPR. Cell Biosci. 2017;7(1):21. doi:10.1186/s13578-017-0148-4.
 126. Shen JP, Zhao D, Sasik R, Luebeck J, Birmingham A, Bojorquez-Gomez A, Licon K, Klepper K, Pekin D,

- Beckett AN. Combinatorial CRISPR–Cas9 screens for de novo mapping of genetic interactions. *Nat Methods*. [2017](#);14(6):573–76. doi:[10.1038/nmeth.4225](#).
127. Zsögön A, Čermák T, Naves ER, Notini MM, Edel KH, Weigl S, Freschi L, Voytas DF, Kudla J, Peres LEP. De novo domestication of wild tomato using genome editing. *Nat Biotechnol*. [2018](#);36(12):1211–16. doi:[10.1038/nbt.4272](#).
 128. Li T, Yang X, Yu Y, Si X, Zhai X, Zhang H, Dong W, Gao C, Xu C. Domestication of wild tomato is accelerated by genome editing. *Nat Biotechnol*. [2018](#);36(12):1160–63. doi:[10.1038/nbt.4273](#).
 129. Lemmon ZH, Reem NT, Dalrymple J, Soyk S, Swartwood KE, Rodriguez-Leal D, Van Eck J, Lippman ZB. Rapid improvement of domestication traits in an orphan crop by genome editing. *Nature Plants*. [2018](#);4(10):766–70. doi:[10.1038/s41477-018-0259-x](#).
 130. Cui Y, Xu J, Cheng M, Liao X, Peng S. Review of CRISPR/Cas9 sgRNA design tools. *Interdiscip Sci Comput Life Sci*. [2018](#);10:455–65.
 131. DeHaan L, Christians M, Crain J, Poland J. Development and evolution of an intermediate wheat-grass domestication program. *Sustainability*. [2018](#);10:1499.
 132. Venske E, Dos Santos Rs, Busanello C, Gustafson P, de Oliveira AC. Bread wheat: a role model for plant domestication and breeding. *Hereditas*. [2019](#);156:1–11.
 133. Zhou J, Xin X, He Y, Chen H, Li Q, Tang X, Zhong Z, Deng K, Zheng X, Akher SA. Multiplex QTL editing of grain-related genes improves yield in elite rice varieties. *Plant Cell Rep*. [2019](#);38(4):475–85. doi:[10.1007/s00299-018-2340-3](#).
 134. Takeda S, Matsuoka M. Genetic approaches to crop improvement: responding to environmental and population changes. *Nat Rev Genet*. [2008](#);9:444–57.
 135. Zhang Y, Massel K, Godwin ID, Gao C. Applications and potential of genome editing in crop improvement. *Genome Biol*. [2018](#);19:210.
 136. Huang L, Zhang R, Huang G, Li Y, Melaku G, Zhang S, Chen H, Zhao Y, Zhang J, Zhang Y, et al. Developing superior alleles of yield genes in rice by artificial mutagenesis using the CRISPR/Cas9 system. *Crop J*. [2018](#);6:475–81.
 137. Lin Z, Yan J, Su J, Liu H, Hu C, Li G, Wang F, Lin Y. Novel OsGRAS19 mutant, D26, positively regulates grain shape in rice (*Oryza sativa*). *Funct Plant Biol*. [2019](#);46:857–68.
 138. Casadevall R, Rodriguez RE, Debernardi JM, Palatnik JF, Casati P. Repression of growth regulating factors by the microRNA396 inhibits cell proliferation by UV-B radiation in Arabidopsis leaves. *Plant Cell*. [2013](#);25:3570–83.
 139. Fina J, Casadevall R, Abdelgawad H, Prinsen E, Markakis MN, Beemster GTS, Casati P. UV-B inhibits leaf growth through changes in growth regulating factors and gibberellin levels. *Plant Physiol*. [2017](#);174:1110–26.
 140. Xie Y, Niu B, Long Y, Li G, Tang J, Zhang Y, Ren D, Liu Y, Chen L. Suppression or knockout of SaF/SaM overcomes the Sa-mediated hybrid male sterility in rice. *J Integr Plant Biol*. [2017](#);59:669–79.
 141. Long SP, Marshall-Colon A, Zhu X-G. Meeting the global food demand of the future by engineering crop photosynthesis and yield potential. *Cell*. [2015](#);161:56–66.
 142. Varshney RK, Thudi M, Roorkiwal M, He W, Upadhyaya HD, Yang W, Bajaj P, Cubry P, Rathore A, Jian J. Resequencing of 429 chickpea accessions from 45 countries provides insights into genome diversity, domestication and agronomic traits. *Nat Genet*. [2019](#);51:857–64.
 143. Cohn M, Bart RS, Shybut M, Dahlbeck D, Gomez M, Morbitzer R, Hou B-H, Frommer WB, Lahaye T, Staskawicz BJ. Xanthomonas axonopodis virulence is promoted by a transcription activator-like effector-mediated induction of a SWEET sugar transporter in cassava. *Mol Plant-Microbe Interact*. [2014](#);27:1186–98.
 144. Xu Z, Xu X, Gong Q, Li Z, Li Y, Wang S, Yang Y, Ma W, Liu L, Zhu B. Engineering broad-spectrum bacterial blight resistance by simultaneously disrupting variable TALE-binding elements of multiple susceptibility genes in rice. *Mol Plant*. [2019](#);12:1434–46.
 145. Oliva R, Ji C, Atienza-Grande G, Huguet-Tapia JC, Perez-Quintero A, Li T, Eom J-S, Li C, Nguyen H, Liu B. Broad-spectrum resistance to bacterial blight in rice using genome editing. *Nat Biotechnol*. [2019](#);37:1344–50.
 146. Liu D, Chen X, Liu J, Ye J, Guo Z. The rice ERF transcription factor OsERF922 negatively regulates resistance to *Magnaporthe oryzae* and salt tolerance. *J Exp Bot*. [2012](#);63:3899–911.
 147. Wang Y, Cheng X, Shan Q, Zhang Y, Liu J, Gao C, Qiu J-L. Simultaneous editing of three homoeoalleles in hexaploid bread wheat confers heritable resistance to powdery mildew. *Nat Biotechnol*. [2014](#);32:947–51.
 148. Nekrasov V, Wang C, Win J, Lanz C, Weigel D, Kamoun S. Rapid generation of a transgene-free powdery mildew resistant tomato by genome deletion. *Sci Rep*. [2017](#);7:1–6.
 149. Langner T, Kamoun S, Belhaj K. CRISPR crops: plant genome editing toward disease resistance. *Annu Rev Phytopathol*. [2018](#);56:479–512.
 150. Arora L, Narula A. Gene editing and crop improvement using CRISPR-Cas9 system. *Front Plant Sci*. [2017](#);8:1932.
 151. Zhenwei Z. The bean seed storage protein β -phaseolin is synthesized, processed, and accumulated in vacuolar type-II protein bodies of transgenic rice endosperm. *Plant Physiol*. [1995](#);109:777–86.
 152. Sindhu AS, Zheng Z, Murai N. The pea seed storage protein legumin was synthesized, processed, and accumulated stably in transgenic rice endosperm. *Plant Sci*. [1997](#);130:189–96.

153. Katsube T, Kurisaka N, Ogawa M, Maruyama N, Ohtsuka R, Utsumi S, Takaiwa F. Accumulation of soybean glycinin and its assembly with the glutelins in rice. *Plant Physiol.* [1999](#);120:1063–74.
154. Burkhardt PK, Beyer P, Wünn J, Klöti A, Armstrong GA, Schledz M, Von Lintig J, Potrykus I. Transgenic rice (*Oryza sativa*) endosperm expressing daffodil (*Narcissus pseudonarcissus*) phytoene synthase accumulates phytoene, a key intermediate of provitamin A biosynthesis. *Plant J.* [1997](#);11:1071–78.
155. FAO. How to Feed the World in 2050. Insights from an Expert Meet FAO [2009](#);2050:1–35.
156. UNICEF [2017](#). World Bank Joint child malnutrition estimates-2018. Geneva, Switzerland: WHO World Bank.