# Induced mutagenesis: an underutilised component in the integrated management of aphid pests in Sub-Saharan Africa

by Zimba, K.J., Sohati, P.H., Munyinda, K., Roberts, J.M. and Pope, T.W.

**Copyright, publisher and additional Information:** This is the author accepted manuscript. The final published version (version of record) is available online via Elsevier. This version is made available under the <u>CC-BY-ND-NC licence</u>

Please refer to any applicable terms of use of the publisher

DOI link to the version of record on the publisher's website



Zimba, K.J., Sohati, P.H., Munyinda, K., Roberts, J.M. and Pope, T.W. (2022) 'Induced mutagenesis: an underutilised component in the integrated management of aphid pests in Sub-Saharan Africa', *Crop Protection*, 159 (106030)

1	Induced mutagenesis: An underutilised component in the integrated management of aphid pests in Sub-Saharan		
2	Africa		
3			
0			
4	Kennedy J. Zimba <sup>a,b*</sup> , Philemon H. Sohati <sup>b</sup> , Kalaluka Munyinda <sup>b</sup> , Joe M. Roberts <sup>a</sup> , Tom W. Pope <sup>a</sup>		
5			
6	<sup>a</sup> Centre for Integrated Pest Management, Agriculture and Environment Department, Harper Adams University,		
7	Newport, Shropshire, TF10 8NB, United Kingdom.		
8	<sup>b</sup> School of Agricultural Sciences, Department of Plant Sciences, University of Zambia, Great East Road Campus,		
9	Lusaka 10101, Zambia.		
10			
11	*Correspondence: zimbaki@gmail.com		
11			
12	Abstract		
13	Aphids (Hemiptera: Aphididae) are important agricultural pests in sub-Saharan Africa. These pests are		
14	primarily controlled by the use of synthetic insecticides, which has consequently led to the emergence of insecticide-		
15	resistant aphid populations as well as negative impacts on non-target organisms. Resistant crop varieties offer a		
16	sustainable approach to manage aphids. Despite regions of sub-Saharan Africa suffering greater crop losses due to		
17	pests, there is only limited availability of genetic engineering and other modern plant breeding technologies. Here		
18	we consider whether induced mutagenesis can contribute to the sustainable management of aphid pests or whether		
19	the lack of research in this area reflects the limitations of this approach.		
20	Keywords: genetic variation, genetic improvement, mutant, plant breeding, resistant varieties		
21			
22	1.0 Introduction		
23	Nearly one billion people are food insecure in Sub-Saharan Africa today (Botha et al., 2020). Predictions		
24	indicate that in the absence of effective mitigation measures, reliable access to sufficient, affordable and nutritious		
25	food is likely to deteriorate further in the next 50 years. Global crop loss estimates per crop of 21.5, 30.0, 22.6, 17.2		
26	and 21.4 % are caused by pests and diseases on wheat, rice, maize, potato and soybean respectively (Savary et al.,		
27	2019). However, these overall estimates mask the very large differences in crop losses among different food		
28	insecurity hotspots such as Sub-Saharan Africa. Aphids (Hemiptera: Aphididae) are amongst the most damaging		

30 crop pests, predominantly relied on the application of synthetic chemical insecticides (Roubos et al., 2014). While

29

invertebrate pests affecting crop productivity (Leybourne et al., 2019). Aphid management has, like for many other

use of such products has greatly enhanced crop productivity during the last century (Naik et al., 2019), issues of resistance due to their overuse as well as their negative effects on human and environmental health are now widely recognised (Kim et al., 2017). Public concerns regarding pesticide use, particularly chemical residues on harvested crops and their products, have also increased significantly during the past few decades (Schaub et al., 2020). These issues have led to increasingly restrictive legislation on insecticide use and, consequently, a diminishing portfolio of effective active ingredients available to growers for invertebrate pest management.

37 Resistant crop varieties offer an environmentally sustainable management option for aphids (Pertot et al., 38 2017). The rapid emergence of plant resistance-breaking aphid biotypes, however, necessitates a regular supply of 39 varieties with new forms of resistance. Despite regions of Sub-Saharan Africa suffering greater crop losses due to 40 pests, availability of genetic engineering and other modern plant breeding technologies (i.e., targeted mutagenesis) 41 are less available (Botha et al., 2020), likely due to the lagging pace in technology, inadequate research funding 42 schemes as well as hesitance of policymakers to establish biosafety laws (Agbowuro et al., 2021; Botha et al., 2020). 43 By contrast, induced mutagenesis is a cost effective, widely accepted tool used for generating genetic variation to 44 abiotic (i.e., drought tolerance) and biotic (i.e., pest resistance) stresses (Singh et al., 2006).

45 Mutagenesis refers to heritable alterations in the genetic material that gives rise to individuals with 46 modified phenotypic traits and provides a source of unique germplasm to facilitate crop improvement (Box 1). Such 47 genetic alterations can be induced by exposing a plant or its propagules to physical or chemical materials with 48 mutagenic properties (Viana et al., 2019). Mutations in the DNA are described based on the alteration of gene 49 functions (Mba, 2013). The common types of mutation induced in the DNA that are relevant to crop improvement 50 include: single base substitutions, point mutations, insertions and deletions (Mba, 2013) (Table 1). Induced 51 mutagenesis has played a key role in the genetic improvement of crops for decades, with the joint Food and 52 Agriculture Organization (FAO) and the International Atomic Energy Agency (IAEA) database containing 3,275 mutant 53 crop varieties derived from 225 plant species (FAO/IAEA, 2019). Mutation derived varieties are now cultivated in 54 most parts of the world, including: Asia, Europe and North America (Horn et al., 2015). Use of induced mutagenesis 55 in plant breeding programs has increased in recent years due to the development of efficient and cost effective 56 mutation-detection techniques such as Targeting Induced Local Lesions in Genomes (TILLING) (Viana et al., 2019). 57 As a technique, induced mutagenesis has been widely adopted by plant breeders targeting pathogen resistance and 58 other abiotic stresses (Oladosu et al., 2016). Few studies, however, have considered using this approach to develop 59 aphid resistant plant varieties. Reasons for why induced mutagenesis should be considered as a means of developing 60 aphid resistant crops in Sub-Saharan Africa are here classified into the following topics: (1) those related to 61 technological issues (i.e., accessibility and legislation) associated with other accelerated approaches to plant 62 breeding, (2) similarities between aphid and pathogen resistance mechanisms, (3) aphid and pathogen resistant genes often being found close together on chromosomes, and (4) improved screening of mutagenised plant 63 64 population.

65

#### 2.0 Prospects for the use of induced mutagenesis as a tool for developing aphid resistant crop varieties

66

## 2.1 Technological issues associated with other accelerated approaches to plant breeding

67 Several crop improvement technologies such as genetic engineering, marker assisted selection and targeted 68 mutagenesis have been developed and may help to accelerate plant breeding for aphid resistance (Bhattacharya, 69 2019; Voss-Fels et al., 2019; Wang et al., 2019). Despite the potential that these modern plant breeding tools offer 70 to plant breeding, their practical use in Sub-Saharan Africa is limited. The use of external DNA in genetically 71 engineered crops, for example, has led to strict biosafety regulation for their use in most Sub-Saharan African 72 countries (Zaidi et al., 2019). In countries like Kenya, for example, where use of genetically engineered crops has 73 been approved, environmental exposure as well as trade of these crops and their products is still prohibited (Botha 74 et al., 2020). Sub-Saharan African countries are largely dominated by smallholder farmers with less financial capacity 75 to annually purchase genetically engineered crop seed (Fischer et al., 2015). There have also been concerns regarding 76 perceived potential risks of genetically enginnered crops on domestic agricultural biodiversity (Jacobsen et al., 2013). 77 In contrast to genetic engineering, targeted mutagenesis involves alteration of endogenous genes (Arora and Narula, 78 2017). Despite the non-integration of external DNA, there is increasing pressure to subject gene-edited crops to the 79 same regulations as crops that are genetically engineered, perhaps due to uncertainty around the intended effects 80 of artificially manipulating plants in this way (Callaway, 2018). In addition, the costs associated with new genomic 81 tools, lack of skilled scientific personnel and laboratories hinder the use of modern molecular approaches to plant 82 breeding in Sub-Saharan Africa (Botha et al., 2020). In comparison to these modern plant breeding tools, induced 83 mutagenesis is more widely used and accepted as a breeding tool with a long history of safe use. The non-84 involvement or use of external DNA in induced mutagenesis exempts mutation derived plants from the often 85 expensive and long regulatory procedures that genetically engineered plants are subjected to (Mba, 2013). This 86 simplified regulatory regime for release of mutant varieties coupled with the robustness, simplicity and low 87 operation costs make induced mutagenesis especially suitable for countries in Sub-Saharan Africa (Mba, 2013).

88

89

## 2.2 Similarities between aphid and pathogen resistance mechanisms

90 Based on the partial overlap between plant-resistance mechanisms against aphids and microbial pathogens 91 (Kaloshian and Walling, 2005), the production of genetic material with disease resistance provides hope for 92 developing aphid resistant cultivars through induced mutagenesis. Plants recognise pathogen-effector proteins (e.g., 93 flagellin, peptidoglycan, lipopolysaccharides in bacteria and chitins in fungi) using receptors on cell walls that trigger 94 defence responses known as pathogen associated molecular pattern (PAMP)-triggered immunity (PTI) (Chisholm et 95 al., 2006). Pathogens, however, have evolved effector proteins that can suppress PTI in plants (Louis et al., 2012). In 96 response, plants have equally evolved additional R proteins that can recognise these pathogen effectors leading to 97 effector-triggered immunity (ETI) (Chisholm et al., 2006). Plants perceive and recognize aphids by detecting specific

98 effector proteins in aphid saliva (e.g., pectinases, cellulases) in a similar way to the detection of pathogens (Dogimont 99 et al., 2010). During feeding, apids inject watery saliva containing proteins and other metabolites into sieve elements 100 (Louis et al., 2012). The protein molecules in aphid saliva are similar to pathogen associated molecular patterns 101 (PAMPs) that are recognised by pattern recognition receptors (PRRs) in plants to trigger PTI (Rodriguez and Bos, 102 2013). To counteract PTI, aphids deliver effector proteins in their host plant to suppress this defense promoting 103 effector-triggered susceptibility (ETS) (Jaouannet et al., 2014). In return, some plant species may carry receptors or 104 R proteins that can recognise effectors in aphid saliva leading to ETI in plants (Jaouannet et al., 2014). Detection of 105 pathogens or aphid species both result in activation of the salicylic acid (SA) signalling pathway likely due to the 106 limited physical damaged to foliage during feeding (Züst and Agrawal, 2016). Indeed, there is evidence that aphids are negatively affected by the activation of the SA pathway. For example, mutant genotypes of Arabidopsis thaliana 107 108 (L.) Heynh with increased SA signalling have been shown to be less susceptible to peach-potato aphid (Myzus 109 persicae Sulzer) (Kerchev et al., 2013). Similarly, growth rate and population growth of the potato aphid 110 (Macrosiphum euphorbiae) is adversely affected by the SA signaling pathway mediated by the Mi-1 gene in tomato 111 (Li et al., 2006). Therefore, it is likely that disease resistant mutants could also resist species of aphid that are 112 vulnerable to the SA signalling pathway.

113

114

## 2.3 Aphid and pathogen resistant genes often being found close together on chromosomes

115 Aphid and pathogen resistance genes are often clustered on the same region of the chromosomes 116 (Dogimont et al., 2010; Seah et al., 2007; Stewart et al., 2009). In apple (Malus domestica Borkh), for example, woolly 117 apple aphid (Eriosoma lanigerum) resistance genes (Er1 and Er2), on chromosomes 8 and 17 respectively, are 118 located on the same genomic regions with genes for resistance to powdery mildew (Bus et al., 2008). The Ra gene 119 on chromosome 2 that mediates resistance in lettuce against the lettuce root aphid (Pemphigus bursarius L.) is 120 clustered together with downy mildew resistance genes on the same chromosome (Christopoulou et al., 2015). 121 Similarly, the potato aphid (*M. euphorbiae*) resistance gene (*Mi*-1) on chromosome 6 in tomato shares the same 122 location (chromosomal region) with disease resistance genes (Seah et al., 2007). Due to this common genomic locale 123 of aphid and pathogen resistance genes, chromosomal alterations due to induced mutagenesis are likely to induce 124 genetic variations for both pathogen and aphid resistance traits.

125

126

#### 2.4 Screening of mutagenised plant population

127 Induced mutagenesis often introduces random changes in the target organism's genome, making it difficult 128 to precisely target specific genes controlling a desired trait. This lack of specificity requires labour intensive screening 129 of large mutant populations (approximately 5,000 to 10,000 genotypes) to optimise chances of finding desirable 130 mutations. To overcome this limitation, techniques such as TILLING (Targeted Induced Local Lesions IN Genomes) 131 have been developed to enhance the detection of useful mutations in mutagenised plant populations (Penna and 132 Jain, 2017). The TILLING technique combines mutagenesis and polymerase chain reaction (PCR) technology to 133 identify point mutations such as single nucleotide polymorphisms (SNPs) in target genes (Irshad et al., 2020). In 134 particular, TILLING allows for identification of variations in mutant genome providing a criteria for shortlisting 135 mutants with potential aphid resistance to include in phenotypic screening (Viana et al., 2019). This molecular 136 approach for identifying mutations, as opposed to whole plants in conventional screening, makes TILLING a high 137 throughput and cost effective screening method. The improved capability of genomic tools in recent years offers 138 more thorough investigations of gene structure and function in mutant genotypes which could allow for easier 139 identification, introgression and molecular characterisation of durable resistance to aphid pests.

140

## 141 **3.0** Discussion and conclusion

142 Relatively few studies have considered the application of induced mutagenesis to develop aphid resistant 143 cultivars, perhaps because induced mutagenesis may result in loss-of-gene function and produces alleles that are 144 often recessive to wild type plants (Sikora et al., 2011). Additionally, induced mutagenesis may alter only one or a 145 few genes producing only minor changes in amino acid composition. Since durable aphid resistance in crops is often 146 mediated by polygenic dominant alleles (Smith and Chuang, 2014), creating polygenic resistance, therefore, is rare 147 using induced mutagenesis (Mba, 2013). However, there have been some notable success in breeding for aphid 148 resistance. Using induced mutagenesis (y-irradiation) on banana (cv. Lakatan), Cueva et al. (2014) succeeded in 149 developing mutants that were repellent and resistant to colonisation by banana aphid (Pentalonia nigronervosa 150 Coquerel). Similarly, Pathak (1991) successfully developed cowpea mutants that were not only repellent but also 151 inhibited survival and reproduction of the cowpea aphid (Aphis craccivora Koch). Mutants derived from turnip 152 cultivars were found to be resistant to mustard aphid (Lipaphis erysimi Kaltenbach), and this resistance was 153 attributed to the non-waxy leaves on these plants. Using a chemical mutagen, Susrama and Pradnyawathi (2019) 154 succeeded in developing mutants of common bean, cowpea and yardlong bean that showed resistance to the 155 cowpea aphid. Similarly, Zimba et al. (in press) showed that mutant cowpea genotypes developed using gamma 156 irradiation reduced colonisation, feeding and population growth of cowpea aphid. Characterisation of feeding 157 behaviour using electrical penetration graph recording indicated that resistance to cowpea aphid in cowpea mutants 158 was mediated by epidermal and mesophyll-based resistance factors.

Use of induced mutagenesis is, however, associated with several limitations. Treatment of plant material by mutagens invariably kills cells causing a wide range of deformities and other side effects (e.g., sterility) in surviving plants (Mba et al., 2010). These deformities are often inherited even in mutant plants with desirable characteristics (Mba et al., 2010). Potential mutants, therefore, usually require several generations of successive propagations or crossing with other genotypes to exclude undesirable side effects from their genetic background (Mba et al., 2010).

5

Mutations arising from induced mutagenesis also have random non-target effects in the genome, making it difficult to precisely target specific genes controlling a desired characteristic (Chaudhary et al., 2019). Therefore, induced mutagenesis programmes are usually 'trial and error' undertakings in which finding a mutant genotype with desirable characteristics is not guaranteed (Chaudhary et al., 2019).

168 The small number of previously reported successes of using induced mutagenesis to produce genotypes 169 with resistance to aphid pests indicate the potential of this approach. Despite this, the far larger number of successes 170 in breeding for disease resistant crops using induced mutagenesis (Busungu et al., 2016; Jung et al., 2005; Oladosu 171 et al., 2016) and the overlap between aphid and pathogen resistance mechanisms indicate that breeding for aphid 172 resistance is a comparatively under exploited use for this technique. This conclusion is further supported by the 173 common location of pathogen and aphid resistant genes on chromosomes. Furthermore, the long history of safe 174 use, low cost of equipment as well as wide acceptability makes induced mutagenesis an important technique that 175 could be exploited further to speed up the delivery of aphid resistant crop varieties in Sub-Saharan Africa. This is 176 emphasised by a current lack of policy frameworks to regulate the use of modern breeding tools in most countries 177 of Sub-Saharan Africa. Although developing aphid resistance using induced mutagenesis is associated with several 178 challenges, this approach provides a practical means through which to develop sustainable management 179 programmes for aphid pests in crops throughout regions such as Sub-Saharan Africa.

180

# 181 Acknowledgement

- 182 Authors are grateful to the International Atomic Energy Agency (IAEA) for the financial support through a sandwich
- 183 PhD fellowship awarded to Kennedy J. Zimba (Grant number: RAF0052-1805114).

184

# References

- Agbowuro, G., Salami, A., Afolabi, M., 2021. Plant breeding: A potential tool to sustain food security in Sub-Saharan Africa. J. Pure Appl. Agric. 6, 1–6.
- Arora, L., Narula, A., 2017. Gene editing and crop improvement using CRISPR-cas9 system. Front. Plant Sci. 8, 1932. https://doi.org/10.3389/fpls.2017.01932
- Bhattacharya, S., 2019. Brassica-aphid interaction: Challenges and prospects of genetic engineering for integrated aphid management. Physiol. Mol. Plant Pathol. 108, 101442. https://doi.org/10.1016/j.pmpp.2019.101442
- Botha, A.M., Kunert, K.J., Maling'a, J., Foyer, C.H., 2020. Defining biotechnological solutions for insect control in sub-Saharan Africa. Food Energy Secur. 9, 1–21. https://doi.org/10.1002/fes3.191
- Bus, V.G.M., Chagné, D., Bassett, H.C.M., Bowatte, D., Calenge, F., Celton, J.M., Durel, C.E., Malone, M.T., Patocchi,
   A., Ranatunga, A.C., Rikkerink, E.H.A., Tustin, D.S., Zhou, J., Gardiner, S.E., 2008. Genome mapping of three
   major resistance genes to woolly apple aphid (Eriosoma lanigerum Hausm.). Tree Genet. Genomes 4, 223–
   236. https://doi.org/10.1007/s11295-007-0103-3
- Busungu, C., Taura, S., Sakagami, J.I., Ichitani, K., 2016. Identification and linkage analysis of a new rice bacterial blight resistance gene from XM14, a mutant line from IR24. Breed. Sci. 66, 636–645. https://doi.org/10.1270/jsbbs.16062

Callaway, E., 2018. CRISPR plants now subject to tough GM laws in European Union. Nature 560(7716), 16-17.

- Chaudhary, J., Alisha, A., Bhatt, V., Chandanshive, S., Kumar, N., Mir, Z., Kumar, A., Yadav, S.K., Shivaraj, S.M.,
  Sonah, H., Deshmukh, R., 2019. Mutation Breeding in Tomato: Advances, Applicability and Challenges. Plants
  8, 1–17. https://doi.org/10.3390/plants8050128
- Chern, M., Xu, Q., Bart, R.S., Bai, W., Ruan, D., Sze-To, W.H., Canlas, P.E., Jain, R., Chen, X., Ronald, P.C., 2016. A Genetic Screen Identifies a Requirement for Cysteine-Rich–Receptor-Like Kinases in Rice NH1 (OsNPR1)-Mediated Immunity. PLoS Genet. 12, 1–20. https://doi.org/10.1371/journal.pgen.1006049
- Chisholm, S.T., Coaker, G., Day, B., Staskawicz, B.J., 2006. Host-microbe interactions: Shaping the evolution of the plant immune response. Cell 124, 803–814. https://doi.org/10.1016/j.cell.2006.02.008
- Christopoulou, M., McHale, L.K., Kozik, A., Wo, S.R.C., Wroblewski, T., Michelmore, R.W., 2015. Dissection of two complex clusters of resistance genes in lettuce (Lactuca sativa). Mol. Plant-Microbe Interact. 28, 751–765. https://doi.org/10.1094/MPMI-06-14-0175-R
- Cueva, F.M., Sison, M.L.J., Dinglasan, E.G., Damasco, O.P., 2014. Elucidating the resistance response of irradiated banana cv. Lakatan to Banana bunchy top virus (BBTV) infection transmitted by the banana aphid Pentalonia

nigronervosa Cocquerel. Philipp. Entomol. 28, 140–154.

- Dogimont, C., Bendahmane, A., Chovelon, V., Boissot, N., 2010. Host plant resistance to aphids in cultivated crops: Genetic and molecular bases, and interactions with aphid populations. Comptes Rendus - Biol. 333, 566–573. https://doi.org/10.1016/j.crvi.2010.04.003
- FAO/IAEA, 2019. The Food and Agriculture Organization of the United Nations/International Atomic Energy Agency – Mutant Variety Database (FAO/ IAEA-MVD) data (2019) reports.
- Feldman, A.B., Leung, H., Baraoidan, M., Elmido-Mabilangan, A., Canicosa, I., Quick, W.P., Sheehy, J., Murchie, E.H., 2017. Increasing leaf vein density via mutagenesis in rice results in an enhanced rate of photosynthesis, smaller cell sizes and can reduce interveinal mesophyll cell number. Front. Plant Sci. 8, 1–10. https://doi.org/10.3389/fpls.2017.01883
- Fischer, K., Van Den Berg, J., Mutengwa, C., 2015. Is Bt maize effective in improving South African smallholder agriculture? S. Afr. J. Sci. 111, 1–2. https://doi.org/10.17159/sajs.2015/a0092
- Guo, H., Mendrikahy, J.N., Xie, L., Deng, J., Lu, Z., Wu, J., Li, X., Shahid, M.Q., Liu, X., 2017. Transcriptome analysis of neo-tetraploid rice reveals specific differential gene expressions associated with fertility and heterosis. Sci.
   Rep. 7, 1–11. https://doi.org/10.1038/srep40139
- Horn, L., Shimelis, H., Laing, M., 2015. Participatory appraisal of production constraints , preferred traits and farming system of cowpea in the northern Namibia : implications for breeding. Legume Research 38, 691– 700. https://doi.org/10.18805/lr.v38i5.5952
- Hussain, A.J., Ali, J., Siddiq, E.A., Gupta, V.S., Reddy, U.K., Ranjekar, P.K., 2012. Mapping of tms8 gene for temperature-sensitive genic male sterility (TGMS) in rice (Oryza sativa L.). Plant Breed. 131, 42–47. https://doi.org/10.1111/j.1439-0523.2011.01897.x
- Hwang, J.E., Ahn, J.W., Kwon, S.J., Kim, J.B., Kim, S.H., Kang, S.Y., Kim, D.S., 2014. Selection and molecular characterization of a high tocopherol accumulation rice mutant line induced by gamma irradiation. Mol. Biol. Rep. 41, 7671–7681. https://doi.org/10.1007/s11033-014-3660-1
- Irshad, A., Guo, H., Zhang, S., Liu, L., 2020. TILLING in cereal crops for allele expansion and mutation detection by using modern sequencing technologies. Agronomy 10 (3), 405. https://doi.org/10.3390/agronomy10030405
- Ishikawa, S., Ishimaru, Y., Igura, M., Kuramata, M., Abe, T., Senoura, T., Hase, Y., Arao, T., Nishizawa, N.K., Nakanishi, H., 2012. Ion-beam irradiation, gene identification, and marker-assisted breeding in the development of low-cadmium rice. Proc. Natl. Acad. Sci. 109, 19166–19171. https://doi.org/10.1073/pnas.1211132109

- Jacobsen, S.E., Sørensen, M., Pedersen, S.M., Weiner, J., 2013. Feeding the world: Genetically modified crops versus agricultural biodiversity. Agron. Sustain. Dev. 33, 651–662. https://doi.org/10.1007/s13593-013-0138-9
- Jaouannet, M., Rodriguez, P.A., Thorpe, P., Lenoir, C.J.G., Macleod, R., Escudero-Martinez, C., Bos, J.I.B., 2014. Plant immunity in plant-aphid interactions. Front. Plant Sci. 5, 1–10. https://doi.org/10.3389/fpls.2014.00663
- Jeng, T.L., Lin, Y.W., Wang, C.S., Sung, J.M., 2012. Comparisons and selection of rice mutants with high iron and zinc contents in their polished grains that were mutated from the indica type cultivar IR64. J. Food Compos. Anal. 28, 149–154. https://doi.org/10.1016/j.jfca.2012.08.008
- Jung, Y.H., Lee, J.H., Agrawal, G.K., Rakwal, R., Kim, J.A., Shim, J.K., Lee, S.K., Jeon, J.S., Koh, H.J., Lee, Y.H., Iwahashi,
   H., Jwa, N.S., 2005. The rice (Oryza sativa) Blast Lesion Mimic Mutant, blm, may confer resistance to blast
   pathogens by triggering multiple defense-associated signaling pathways. Plant Physiol. Biochem. 43, 397–
   406. https://doi.org/10.1016/j.plaphy.2005.03.002
- Kaloshian, I., Walling, L.L., 2005. Hemipterans as plant pathogens. Annu. Rev. Phytopathol. 43, 491–521. https://doi.org/10.1146/annurev.phyto.43.040204.135944
- Kerchev, P.I., Karpińska, B., Morris, J.A., Hussain, A., Verrall, S.R., Hedley, P.E., Fenton, B., Foyer, C.H., Hancock,
   R.D., 2013. Vitamin C and the abscisic acid-insensitive 4 transcription factor are important determinants of aphid resistance in arabidopsis. Antioxidants Redox Signal. 18, 2091–2105.
   https://doi.org/10.1089/ars.2012.5097
- Kim, B., Woo, S., Kim, M.J., Kwon, S.W., Lee, J., Sung, S.H., Koh, H.J., 2018. Identification and quantification of flavonoids in yellow grain mutant of rice (Oryza sativa L.). Food Chem. 241, 154–162. https://doi.org/10.1016/j.foodchem.2017.08.089
- Kim, K.H., Kabir, E., Jahan, S.A., 2017. Exposure to pesticides and the associated human health effects. Sci. Total Environ. 575, 525–535. https://doi.org/10.1016/j.scitotenv.2016.09.009
- Leybourne, D.J., Valentine, T.A., Robertson, J.A.H., Pérez-Fernández, E., Main, A.M., Karley, A.J., Bos, J.I.B., 2019. Defence gene expression and phloem quality contribute to mesophyll and phloem resistance to aphids in wild barley. J. Exp. Bot. 70, 4011–4026. https://doi.org/10.1093/jxb/erz163
- Li, Q., Xie, Q.G., Smith-Becker, J., Navarre, D.A., Kaloshian, I., 2006. Mi-1-mediated aphid resistance involves salicylic acid and mitogen-activated protein kinase signaling cascades. Mol. Plant-Microbe Interact. 19, 655– 664. https://doi.org/10.1094/MPMI-19-0655
- Lin, D.G., Chou, S.Y., Wang, A.Z., Wang, Y.W., Kuo, S.M., Lai, C.C., Chen, L.J., Wang, C.S., 2014. A proteomic study of rice cultivar TNG67 and its high aroma mutant SA0420. Plant Sci. 214, 20–28.

https://doi.org/10.1016/j.plantsci.2013.09.010

- Long, W., Dong, B., Wang, Yihua, Pan, P., Wang, Yunlong, Liu, L., Chen, X., Liu, X., Liu, S., Tian, Y., Chen, L., Wan, J., 2017. FLOURY ENDOSPERM8, encoding the UDP-glucose pyrophosphorylase 1, affects the synthesis and structure of starch in rice endosperm. J. Plant Biol. 60, 513–522. https://doi.org/10.1007/s12374-017-0066-3
- Louis, J., Singh, V., Shah, J., 2012. Arabidopsis thaliana Aphid Interaction . Arab. B. 10, e0159. https://doi.org/10.1199/tab.0159
- Mba, C., 2013. Induced Mutations Unleash the Potentials of Plant Genetic Resources for Food and Agriculture. Agronomy 3, 200–231. https://doi.org/10.3390/agronomy3010200
- Mba, C., Afza, R., Bado, S., Jain, S.M., 2010. Induced mutagenesis in plants using physical and chemical agents, in: Davey, M.R., Anthony, P. (Eds.), Plant Cell Culture: Essential Methods. John Wiley & Sons, London. pp. 111– 130.
- Naik, K., Mishra, S., Srichandan, H., Singh, P.K., Sarangi, P.K., 2019. Plant growth promoting microbes: Potential link to sustainable agriculture and environment. Biocatal. Agric. Biotechnol. 21, 101326.
   https://doi.org/10.1016/j.bcab.2019.101326
- Oladosu, Y., Rafii, M.Y., Abdullah, N., Hussin, G., Ramli, A., Rahim, H.A., Miah, G., Usman, M., 2016. Principle and application of plant mutagenesis in crop improvement: A review. Biotechnol. Biotechnol. Equip. 30, 1–16. https://doi.org/10.1080/13102818.2015.1087333
- Pathak, R.S., 1991. Genetic evaluation of two aphid resistant cowpea mutants in Kenya, in: Wang, L., (Eds.), Plant Mutation Breeding for Crop Improvement. inis.iaea.org. Vienna, Austria. pp. 241–246.
- Penna, S., Jain, S.M., 2017. Mutant resources and mutagenomics in crop plants. Emirates J. Food Agric. 29, 651– 657. https://doi.org/10.9755/ejfa.2017.v29.i9.86
- Pertot, I., Caffi, T., Rossi, V., Mugnai, L., Hoffmann, C., Grando, M.S., Gary, C., Lafond, D., Duso, C., Thiery, D.,
   Mazzoni, V., Anfora, G., 2017. A critical review of plant protection tools for reducing pesticide use on
   grapevine and new perspectives for the implementation of IPM in viticulture. Crop Prot. 97, 70–84.
   https://doi.org/10.1016/j.cropro.2016.11.025
- Phanchaisri, B., Chandet, R., Yu, L.D., Vilaithong, T., Jamjod, S., Anuntalabhochai, S., 2007. Low-energy ion beaminduced mutation in Thai jasmine rice (Oryza sativa L. cv. KDML 105). Surf. Coatings Technol. 201, 8024– 8028. https://doi.org/10.1016/j.surfcoat.2006.02.057
- Rodriguez, P.A., Bos, J.I.B., 2013. Toward understanding the role of aphid effectors in plant infestation. Mol. Plant-Microbe Interact. 26, 25–30. https://doi.org/10.1094/MPMI-05-12-0119-FI

- Roubos, C.R., Rodriguez-Saona, C., Isaacs, R., 2014. Mitigating the effects of insecticides on arthropod biological control at field and landscape scales. Biol. Control 75, 28–38. https://doi.org/10.1016/j.biocontrol.2014.01.006
- Ruengphayak, S., Ruanjaichon, V., Saensuk, C., Phromphan, S., Tragoonrung, S., Kongkachuichai, R., Vanavichit, A., 2015. Forward screening for seedling tolerance to Fe toxicity reveals a polymorphic mutation in ferric chelate reductase in rice. Rice 8(1), 1-10. https://doi.org/10.1186/s12284-014-0036-z
- Savary, S., Willocquet, L., Pethybridge, S.J., Esker, P., McRoberts, N., Nelson, A., 2019. The global burden of pathogens and pests on major food crops. Nat. Ecol. Evol. 3, 430–439. https://doi.org/10.1038/s41559-018-0793-y
- Schaub, S., Huber, R., Finger, R., 2020. Tracking societal concerns on pesticides A Google Trends analysis. Environ. Res. Lett. 15(8), 084049. https://doi.org/10.1088/1748-9326/ab9af5
- Seah, S., Telleen, A.C., Williamson, V.M., 2007. Introgressed and endogenous Mi-1 gene clusters in tomato differ by complex rearrangements in flanking sequences and show sequence exchange and diversifying selection among homologues. Theor. Appl. Genet. 114, 1289–1302. https://doi.org/10.1007/s00122-007-0519-z
- Sikora, P., Chawade, A., Larsson, M., Olsson, J., Olsson, O., 2011. Mutagenesis as a tool in plant genetics, functional genomics, and breeding. Int. J. Plant Genomics 2011, 1-13. https://doi.org/10.1155/2011/314829
- Singh, V.V., Ramkrishna, K., Arya, R.K., 2006. Induced chemical mutagenesis in cowpea [Vigna unguiculata (L.) Walp.]. Indian J. Genet 66(4), 312–315.
- Smillie, I., Pyke, K., Murchie, E., 2012. Variation in vein density and mesophyll cell architecture in a rice deletion mutant population. J. Exp. Bot. 63, 4563–4570. https://doi.org/10.1093/jxb/err313
- Smith, C.M., Chuang, W., 2014. Plant resistance to aphid feeding : behavioral , physiological , genetic and molecular cues regulate aphid host selection and feeding. Pest Manag. Sci. 70, 528–540. https://doi.org/10.1002/ps.3689
- Song, J.Y., Kim, D.S., Lee, M.C., Lee, K.J., Kim, J.B., Kim, S.H., Ha, B.K., Yun, S.J., Kang, S.Y., 2012. Physiological characterization of gamma-ray induced salt tolerant rice mutants. Aust. J. Crop Sci. 6, 421–429.
- Stewart, S.A., Hodge, S., Ismail, N., Mansfield, J.W., Feys, B.J., Prospéri, J.M., Huguet, T., Ben, C., Gentzbittel, L., Powell, G., 2009. The RAP1 gene confers effective, race-specific resistance to the pea aphid in Medicago truncatula independent of the hypersensitive reaction. Mol. Plant-Microbe Interact. 22, 1645–1655. https://doi.org/10.1094/MPMI-22-12-1645

Susrama, I.G.K., Pradnyawathi, N.L.M., 2019. Induced In Vivo mutagenesis using colchisine in hydrogen peroxide on

M2 cowpea, yardlong bean and common bean. Int. J. Biosci. Biotechnol. 7, 1–11.

- Tu, Y., Jiang, A., Gan, L., Hossain, M., Zhang, Jinming, Peng, B., Xiong, Y., Song, Z., Cai, D., Xu, W., Zhang, Jianhua, He, Y., 2014. Genome duplication improves rice root resistance to salt stress. Rice 7, 1–13. https://doi.org/10.1186/s12284-014-0015-4
- Viana, V.E., Pegoraro, C., Busanello, C., Costa de Oliveira, A., 2019. Mutagenesis in Rice: The Basis for Breeding a New Super Plant. Front. Plant Sci. 10, 1–28. https://doi.org/10.3389/fpls.2019.01326
- Voss-Fels, K.P., Stahl, A., Hickey, L.T., 2019. Q&A: Modern crop breeding for future food security. BMC Biol. 17, 1– 7. https://doi.org/10.1186/s12915-019-0638-4
- Wang, X., Gao, Y., Chen, Z., Li, J., Huang, J., Cao, J., Cui, M., Ban, L., 2019. (Ε)-β-farnesene synthase gene affects aphid behavior in transgenic Medicago sativa. Pest Manag. Sci. 75, 622–631. https://doi.org/10.1002/ps.5153
- Xu, J., Shi, S., Wang, L., Tang, Z., Lv, T., Zhu, X., Ding, X., Wang, Y., Zhao, F.J., Wu, Z., 2017. OsHAC4 is critical for arsenate tolerance and regulates arsenic accumulation in rice. New Phytol. 215, 1090–1101.
   https://doi.org/10.1111/nph.14572
- Zaidi, S.S.E.A., Vanderschuren, H., Qaim, M., Mahfouz, M.M., Kohli, A., Mansoor, S., Tester, M., 2019. New plant breeding technologies for food security. Science (80-. ). 363, 1390–1391. https://doi.org/10.1126/science.aav6316
- Züst, T., Agrawal, A.A., 2016. Mechanisms and evolution of plant resistance to aphids. Nat. Publ. Gr. 1–9. https://doi.org/10.1038/nplants.2015.206

Mutagens	Type of mutation	Mutation derived traits	References		
Chemical agents					
Ethyl Methanesulfonate	Guanine alkylation, G/C to A/T	Plant development and metabolism	(Feldman et al., 2017)		
	transitions or G/C to C/G or G/C to	Abiotic stress tolerance	(Xu et al., 2017)		
	T/A transversions				
N-methyl-N-nitrosourea	Guanine and	Biotic stress tolerance	(Busungu et al., 2016)		
	cytosine alkylation, G/C to T/A	Nutritional quality	(Kim et al., 2018)		
	transitions	Yield and quality improvement	(Long et al., 2017)		
Sodium azide	Generates azidoalanine	Abiotic stress tolerance	(Hussain et al., 2012)		
	causing G/C to A/T transitions	Nutritional improvement	(Jeng et al., 2012)		
		Yield and quality improvement	(Lin et al., 2014)		
Colchicine	Chromosome doubling, affects the	Nutritional improvement	(Viana et al., 2019)		
	microtubules promoting symmetric	Abiotic stress tolerance	(Tu et al., 2014)		
	cell division.	Yield and quality improvement	(Guo et al., 2017)		
Physical agents					
Gamma-Rays	Single nucleotide substitution,	Plant development and metabolism	(Smillie et al., 2012)		
	inversion and deletion	Abiotic stress tolerance	(Song et al., 2012)		
		Nutritional improvement	(Hwang et al., 2014)		
Ion Beam Radiation	Point mutation (deletion), inversion,	Plant development and metabolism	(Phanchaisri et al., 2007)		
	translocation and insertion	Nutritional quality	(Ishikawa et al., 2012)		
Fast-Neutron Irradiation	A/T to G/C	Abiotic stress tolerance	(Ruengphayak et al., 2015)		
	transition, insertion, inversion,	Biotic stress resistance	(Chern et al., 2016)		
	duplication and deletion				

**Table 1**: Major chemical and physical mutagens used for induction of random mutations in plants

# Box 1. Illustration of a generalised procedure for induced mutagenesis.

Seed is denoted as  $'M_0'$  before mutagen treatment and  $'M_{1.n'}$  for generations following mutagenesis. 'M' = meotic generation. After mutagenesis,  $M_1$  seed is planted to produce  $M_1$  plants and  $M_2$  seed. Due to heterozygosity of  $M_1$  plants, mutations are not yet visible at this stage. Seed is harvested, bulked and planted to produce  $M_2$  plants and  $M_3$  seed. Mutations begin to appear in the  $M_2$  population due to genetic recombination and segregations which marks the beginning of screening and selection of desired mutants. From  $M_3$  onwards, seed is harvested from individual plants and planted as single plant-progenies to facilitate detailed screening of mutants. Several generations (i.e.  $M_{3.6}$ ) are required for mutant genotypes to reach homogeneity. Homogenous mutants (i.e.  $M_{5.6}$ ) with desired traits can be directly used as a variety or as parents in breeding programmes

