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#### Abstract

The floriculture industry in developing countries has grown significantly due to improved production methods and rising demand for unique, resilient flowers. Biotechnological methods like micropropagation and genetic modification are crucial for enhancing ornamental plants, including traits such as floral morphology, color, fragrance, stress resistance, and post-harvest longevity. This review emphasizes the importance of biotechnology in creating commercially desirable varieties and understanding the chemical composition of edible flowers. Edible flowers, with historical significance and health benefits, are gaining renewed interest. Advances in biotechnological methods like mutation breeding, molecular markers, genetic engineering and genome editing offer promise in meeting global demand for nutritious food options through enhanced edible flowers, highlighting the need for ongoing research and innovation.

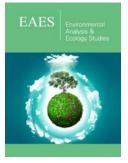
Keywords: Edible flowers, Advance biotechnological tools, Nutritional composition, Genetic engineering, Floriculture industry

### Introduction

Edible flowers possess a rich history of consumption and documentation worldwide, spanning ancient civilizations like Greece, Rome, medieval Europe and Asian countries such as China and Japan [1,2]. Over time, globalization and heightened consumer awareness have rekindled interest in edible flowers due to their potential to enhance human well-being and health. Research has spotlighted their bioactive compounds, including natural pigments, essential oils and antioxidants, elucidating their health-promoting effects and folk medicinal uses. The common phytochemicals in edible flowers are depicted in (Table 1). In response to consumer preferences for natural, functional and healthy food products, edible flowers have gained considerable attraction in the market which led to the evaluation of several species such as chrysanthemum, hibiscus, lavender, marigold and rose for their potential benefits [13,14]. Approximately 180 flower species have been identified as suitable for human consumption, edible flowers offer more than just aesthetic enhancement; they present a safe and nutritious option. These blossoms play a crucial role as functional ingredients in food along with their aroma and have potential health advantages when included in range of dishes and beverages like teas, wines, fruit juices etc. [15,2]. Beyond culinary appeal, their extensive historical use in traditional medicine underscores their medicinal properties.



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Table 1: C	Common	phytochemicals	in	edible	flowers.
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Edible Flowers	Phytochemicals	Reference	
Marigold, Rose, Calendula, Chrysanthemum	Carotenoids (Lutein, $\beta$ -carotene, Flavoxanthin, lycopene, zeaxanthin, etc)	Varzakas and Kiokias [3]; Wan et al. [4]; Pavelkova et al. [5]	
Rose, Hibiscus, Calendula, Chrysanthemum	Anthocyanins (Cyanidin 3,5-diglucoside, pelargonidin 3,5-diglucoside, peonidin 3-O-glucoside, etc)	Grajeda-Iglesias et al. [6]; Wan et al. [7]; Kumari et al. [8]	
Rose, <i>Calendula</i> , Marigold, Lavender, <i>Hibiscus</i> , Daylily, <i>Chrysanthemum</i>	Flavonoids (Quercetin, myricetin 3,5-di-O-glucoside, kaempferol 3,7-di-O-rhamnoside, etc)	Cendrowski et al. [9]; Cao et al. [10]; Wan et al. [4]	
Rose, Lavender, <i>Chrysanthemum, Calendula, Daylily</i> , Marigold	Phenolic acid (Gallic, Caffeic, Caftaric, Chlorogenic, Chicoric, Coumaric, Sinapic and Ferulic, etc)	Ryu et al. [11]; Krzymińska et al. [12]	

As a niche market, edible flowers cater to a diverse range of consumers seeking natural and bioactive components in their diet, further highlighting their significance in contemporary food culture. In the present scenario, there is a global demand for novel traits in ornamental plant products, including improved anatomical attributes, floral colors, pigments, stress tolerance, disease resistance and enhanced secondary metabolites with health benefits and medicinal values [16,17]. Research and advancements in biotechnology Figure 1 have paved the way to enhance edible flowers, ensuring their safety and augmenting their valuable traits for diverse purposes, including industrial applications, promising a future for these blossoms.

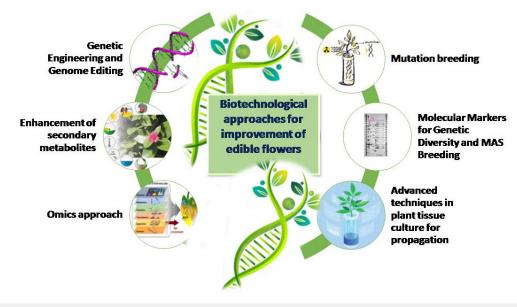


Figure 1: Biotechnological approaches for improvement of edible flowers.

## **Biotechnological Approaches Employed for Improving Specific Traits in Edible Flower Crops**

### **Mutation breeding**

Mutation breeding has been a fundamental approach in developing new plant varieties, especially in ornamental plants. Traditional breeding methods, although reliable, are laborious and time-consuming compared to mutation breeding. This technique involves the use of UV radiation, ionizing radiation and chemical mutagens like EMS and Sodium Azide to induce mutations. It has significantly enhanced breeding efficiency. Mutation breeding revolutionized ornamental plant breeding, accelerating the development of novel traits, particularly flower colors. This method is pivotal in satisfying the floriculture industry's demands for unique attributes. Various techniques, including gamma radiation optimization and chemical mutagen selection, have yielded promising results in plants like carnations, *dianthus*, *chrysanthemums*, roses and more. These methods have shown potential in creating new cultivars and enhancing characteristics like flower color and plant architecture. Research institutions have actively registered number of new mutant lines of *dianthus*, *chrysanthemums*, roses, marigold, Jasmine, tuberose, aster, gladiolus and many more, ensuring the continuous innovation of ornamental plant varieties [18-29] DFR Annual Report, 2022-23). Mutation breeding methods are pivotal for the growth and innovation of the ornamental plant industry, enabling the creation of desirable traits to meet local and export market demands.

# Molecular markers for genetic diversity and MAS breeding

Molecular markers have emerged as valuable tools for genetic improvement in ornamental plants, offering insights into genetic diversity and aiding in various applications such as Marker-Assisted Selection (MAS). Various molecular techniques like SSR, ISSR and SNP have been effectively utilized to assess genetic diversity, determine parentage and identify specific traits. These markers have been employed in genetic studies of different ornamental species, revealing unique alleles and providing essential information for plant breeding program [30]. For example, SSR markers have been used to distinguish different rose varieties and evaluate genetic diversity in chrysanthemums [31,32] Additionally, GWAS studies have linked specific genes to traits like flower type and shape in chrysanthemums [33]. Agarwal et al. [34] identified the polymorphic nature of SCoT markers and established genetic diversity among different Rosa germplasms. Rosa hybrida's genetic fidelity was affirmed with RAPD and ISSR markers [35] and SSR markers differentiated between cultivated and wild rose species [36,37] constructed the first individual maps of rose populations, linking them with 824 SNPs and 13 SSR bridge markers. In carnations, molecular markers unveiled the genetic basis of resistance to Fusarium oxysporum, with a sequence-specific PCRbased SCAR marker developed [38]. SSR markers differentiated carnation varieties [39], while RAPD markers assessed genetic fidelity of in vitro regenerated plants [40]. Marigold research employed molecular markers for investigating male sterility [41] and assessing genetic diversity in genotypes. In Lilium species, ISSR and AFLP markers confirmed genetic stability and variations in regenerants and progeny [42,43]. ICAR-DFR is working on unique fingerprints for marigold, rose and chrysanthemum, employing SRAP, ISSR, and SCoT markers, with a specific focus on varietyspecific markers in chrysanthemums (Unpublished data).

# Advanced techniques in plant tissue culture for propagation

The floriculture industry, driven by the demand for high-quality ornamental plants and flowers, consistently seeks new varieties with improved traits, including plant architecture, vase life, keeping quality, color, and disease resistance [16,17]. Ornamental horticulture encompasses a wide range of plants used in various applications, such as home gardening, landscaping and cut flower production, contributing significantly to the industry [44,45]. Tissue culture technology has brought about a revolution in the floriculture sector, particularly through large-scale propagation using in vitro cloning. Tissue culture has proven invaluable in generating genetic variability, enhancing plant health and expanding germplasms available to breeders. This technology has successfully incorporated specific traits through gene transfer, resulting in new genetic variations within breeding lines [36]. Various in vitro techniques, such as protoplast culture, organ culture and meristem culture, have been employed to produce haploid lines and Somaclonal variation in Table 1, offering advantages over traditional methods in terms of time and mass multiplication [17]. In addition to mass production, tissue culture has enabled pathogen elimination from planting materials and the production of uniform, disease-free plantlets for large-scale cultivation. These techniques have also facilitated the rapid multiplication of chimeric-mutant plants obtained through mutagenesis [16]. In vitro techniques have significantly reduced the time required for developing new varieties and have facilitated the rapid dissemination of planting materials over large areas [46]. (Table 2) has summarized the tissue culture techniques employed for edible flower species. ICAR-DFR has developed efficient regeneration protocols for indigenous chrysanthemum varieties, which are particularly valuable for multiplying chimeric flowers induced by physical mutagens. The institute has also established regeneration and direct organogenesis protocols for marigold varieties, allowing for the development of transgenic and genomeedited plants (Unpublished data).

Table 2: Summary of tissue culture techniques employed for edible flower species.

Species/Cultivars	Explant	Response	References		
Organogenesis					
Chrysanthemum spp.	Shoot tips, Nodal explants and leaf	Multiple shoot, root and plant formation	Jahan et al. 2021		
Gerbera jamesonii	Shoot tips, floral buds, leaf, petioles and petals	Multiple shoot root and plant formation	Winarto B & Prama Yufdy M 2017; Akter et al. 2012		
Hemerocallis fulva	Stem tissue	Callus and shoot formation	Matand et al. 2020		
Hibiscus rosa sinensis	Nodal explants with axillary bud	Multiple shoot root and plant formation	Metwally et al. 2016		
Jasminum sambac	Young leaves, stems and petioles	Callus and shoot formation	Farzinebrahimi et al. 2014		
Dianthus caryophyllus	Axillary buds	Multiple shoot root and plant formation	Ahmadian et al. 2017		
Nelumbo nucifera	Immature cotyledon and embryo and meristem and embryos	Callus induction and Multiple shoo, root and plant formation	Deng et al. 2020 and Yu et al. 2015		
Rhododendron arboretum	Nodal explants	Multiple shoot root and plant formation	Mao et al. 2018		
Rosa spp.	Axillary buds	Multiple shoot root and plant formation	Attia et al. 2012 and Baig et al. 2011		
Tagetes erecta and Tagetes patula	Nodal segments, shoot tip	Multiple shoot root and plant formation	Kumar et al. 2018, Majumder et al. 2014		
Soma clonal variation					
Chrysanthemum spp.	Adventitious shoots from two explant types, leaves and internodes Eeckhau et al. 2020 and Zalewska et al. 2011				

Gerbera	Capitulum	Bhatia et al. (2009)			
Carnation	Leaf base explants	Esmaiel et al. (2012)			
Somatic embryogenesis					
Rosa spp.	Nodal stem segment	Li et al. 2002 and Kim et al. 2003			
	Haploid development				
Chrysanthemum spp.	Anther, Ovule	Khandakar et al. 2014 and Wang et al. 2014 and Gao et al. 2010			
Carnation	Anther	Nontaswatsri et al. 2007			
Lily	Anther	Arzate-Fernandez et al. 1997			
Marigold	Anther	Kurimella et al. 2021			
Gerbera	Ovule	Cappadocia et al. 1988			

# Genetic engineering and genome editing for edible flower crops

Genetic manipulation techniques have emerged as powerful tools to overcome limitations of traditional breeding methods in the ornamental plant industry. Genetic engineering allows the introduction of desired genes into ornamental plants, providing the ability to modify traits such as color, fragrance, disease resistance and flower architecture. *Agrobacterium*-mediated plant genetic transformation is a widely used method for gene transfer due to its simplicity and adaptability (Table 3). Through genetic engineering, it is possible to enhance the quality and traits of ornamental plants for both aesthetic and practical purposes. This approach offers a promising way to develop new cultivars with novel flower colors, improved post-harvest longevity and other desired characteristics, contributing to the advancement of the ornamental plant industry described below.

Table 3: Agrobacterium-mediated stable gene transformation in edible flower plants.

Species	Exogenous	Explant	Methods	Gene A. <i>tumefaciens</i> strain	Transformation Efficiency	Phenotype of Transgenic Plant	Ref
Rosa hybrida	GFP	Leaf	Somatic embryogenesis	GV3101	5~6%	Green fluorescence observed	Liu et al. 2021
Rosa chinensis	GUS	Somatic embryos	Somatic embryogenesis+Shoot regeneration	EHA105	ND	GUS positive	Vergne et al. 2010
Tagetes erecta	GFP	Flower	Floral dipping	EHA105	ND	Green fluorescence observed	Cheng et al., 2019
Tagetes erecta	GUS	Leaf	Shoot regeneration	LBA4404	ND	GUS positive in leaves of transgenic plants	Narushima et al., 2017
Chrysanthemum	Artemisinin biosynthesis genes	Leaf	Shoot regeneration	CBE21	0.17~0.33%	Artemisinin production	Firsov et al., 2020
Chrysanthemum	RsMYB1	Leaf	Shoot regeneration	GV3101	1%	Improved resistance to herbicides	Naing et al., 2016
Chrysanthemum	cry1Ab	Leaf	Shoot regeneration	LBA4404	ND	Improved insect resistance	Shinoyama et al., 2002
Lilium 0-	GUS	Meristematic nodular calli	Shoot regeneration	EHA101	11.10%	Stable expression of GUS gene	Abbasi et al., 2020
Lilium	GUS	Filament- derived calli	Shoot regeneration	EHA101	ND	GUS positive	Hoshi et al., 2004
Gerbera hybrida	GMYB10	Leaf	Shoot regeneration	C58C1	ND	Activation of Anthocyanin Biosynthesis	Elomaa et al., 2003

**Flower colour:** Plant flower colors depend on anthocyanins, water-soluble pigments that bring vibrant hues to fruits, flowers and leaves. These pigments, derived from six anthocyanidins, are pivotal in genetic and molecular research [47]. Anthocyanin creation involves a series of enzymatic reactions within the flavonoid biosynthetic pathway, encompassing phenylalanine,

phenylpropanoid, flavonoid and anthocyanin metabolism. In the realm of genetic manipulation for flower color, *Petunia hybrida* with its well-studied genetics and transformability, stands out as a valuable model for engineering flower color [48]. *Chrysanthemum*, known for its diverse flower colors, has been genetically engineered to intensify cyanidin content for vibrant red petals.

However, achieving blue flower color through the introduction of a pansy F3'5'H gene remains a persistent challenge [49]. Moon carnations and roses have been successfully engineered to display violet and blue flower colors, respectively. Introduction of petunia genes F3'5'H and DFR into white carnation mutants resulted in violet hues through co-pigmentation. Similarly, roses were modified to produce delphinidin-based anthocyanins, leading to blue and magenta flower colors via specific gene introduction [50]. Transient transformation experiments in Lilium plants have demonstrated the alteration of flower color through the overexpression of key genes for instance, overexpressing the Phalaenopsis F3'5'H (Ph F3'5'H) gene shifted the color from pink to pale purple, with further deepening of the purple color achieved by co-expressing PhF3'5'H and Hyacinth DFR (HyDFR) genes [51]. Genome editing has emerged as a potent tool for modifying flower color across various plant species. Understanding anthocyanin biosynthesis and related genes offers exciting possibilities for creating unique flower colors, greatly impacting the ornamental plants industry with visually captivating varieties.

Floral scent: Floral scent plays a vital role in attracting pollinators, protecting plants from herbivores and pathogens and facilitating reproductive processes. The scent is a complex mixture of compounds, including fatty acid derivatives, benzenoids, phenylpropanoids and terpenoids. Understanding the intricate role of floral scent in plant biology offers opportunities to improve crop yields and enhance the visual and olfactory appeal of flowers [52,53]. Genetic engineering offers a solution to impart fragrance to modern cut-flowers, addressing the decline of natural scent due to selective breeding for other traits. A breakthrough involved isolating the S-Linalool Synthase (LIS) gene from Clarkia breweri, a sweet-scented California plant. By introducing the LIS gene into carnations, linalool production was successfully enhanced, showcasing the potential of genetic manipulation to enhance floral scents [54]. Transcription Factors (TFs) are key players in fragrance biosynthesis, regulating various pathways that lead to the production of volatile compounds. They control processes like the shikimate pathway, regulated by TF ODORNT1 (ODO1), specifically expressed in *petunia* petals. While some pathways are well understood, the regulation of terpenoid pathways remains unclear. The successful introduction of the PAP1 transcription factor from Arabidopsis into rose flowers resulted in heightened production of specific scent compounds, emphasizing the potential of genetic engineering to enhance floral fragrances and revolutionize the floral industry [55].

**Biotic and abiotic stress resistance:** Plants are susceptible to a wide range of environmental stresses viz., abiotic (e.g., radiation, salinity, drought) and biotic (e.g., pathogens, herbivores) that impair growth and productivity [17]. These stresses trigger diverse plant responses, altering gene expression, metabolic processes and physiological attributes. Research on stress tolerance in edible flower crops has made progress in understanding and improving their responses, exemplified by transgenic carnations with enhanced resistance to Fusarium wilt disease and caffeineproducing chrysanthemum plants exhibiting resistance to fungal attacks [56,57]. The abundance, organization and expression patterns of LlWRKY genes were studied in Lilium, indicating their potential role in both abiotic and biotic stress tolerance [58]. Genetic transformation methods, like *Agrobacterium*-mediated transfer of specific genes, have been instrumental in conferring resistance against viruses and pests, enhancing crop resilience. Additionally, the role of critical transcription factors such as ZIP, WRKY and NAC in stress response pathways has been demonstrated, paving the way for targeted approaches to enhance stress tolerance in ornamental plants [59,60]. Overall, employing biotechnological strategies holds promise in mitigating environmental stress effects and improving the productivity and stress resilience of ornamental plants.

Keeping quality and post-harvest quality management: Ornamental plants are integral to the horticulture industry and preserving their quality and freshness is commercially important. Ethylene and bacterial infections accelerate flower wilting, necessitating resistance. Genetic engineering offers a cost-effective solution by reducing ethylene-induced senescence [61]. Targeting ethylene-related genes enhances vase life and quality in flowers. Successful genetic modifications, as seen in carnations, illuminate this potential [62]. Additionally, optimizing tissue-specific ethylene insensitivity, demonstrated in transgenic petunias, is crucial for commercial viability. In roses, various genes and transcription factors have been implicated in the ethylene response, providing opportunities to manipulate floral senescence and extend flower life. In conclusion, genetic engineering focused on ethylene pathways holds promise for enhancing the shelf life of ornamental plants, ensuring better quality and commercial value. Understanding regulatory mechanisms and gene expression patterns in ethylene response pathways is pivotal for developing enduring ornamental plants.

Enhancement of industrially important pigments, essential oils and secondary metabolites: Highlighting the nutraceutical potential of edible flowers, underscoring their richness in nutrients, phytochemicals and the subsequent health benefits, these flowers demonstrate a broad range of medicinal attributes, including anti-diabetic, anti-cancer, anti-anxiety, anti-inflammatory, antimicrobial, diuretic and immunomodulatory effects [63,64]. Also, flower essential oils are highly valued for their aromatic appeal and pharmacological properties enriched with active constituents. These oils possess antimicrobial, antioxidant and anti-pest properties, crucial for food preservation, cosmetics and medicine [65,66]. Biotechnological advancements show potential in optimizing essential oil yield and composition, exemplified by studies focusing on lavender as a key example [67-69].

**Omics approaches:** Crop improvement endeavors aim to develop climate-smart crops with heightened stress tolerance, improved nutritional value and superior agronomic traits. The integration of "omics" technologies-genomics, transcriptomics, proteomics and metabolomics-has been pivotal in identifying pivotal genes, proteins and metabolic pathways governing desired traits, facilitating marker-assisted breeding in major crops. Additionally, harnessing natural variation in crop wild relatives and underutilized species is crucial.

#### Conclusion

Edible flowers are highly valued for their functional properties, adding appeal and nutritional benefits to food products. With growing consumer demand for natural and healthy food options, there's a significant opportunity to use biotechnological tools to enhance the quality and yield of edible flowers to meet global demands. Progress has been notable in enhancing key traits through mutation breeding and tissue culture techniques. Despite genome sequencing progress, the application of genome editing in edible ornamentals is still evolving. There's a need to extend transgenic, genome editing and omics technologies to enhance the nutritional and bioactive properties of valuable edible flowers.

#### References

- 1. He J, Yin T, Chen Y, Cai L, Tai Z, et al. (2015) Phenolic compounds and antioxidant activities of edible flowers of *pyrus pashia*. Journal of Functional Foods 17: 371-379.
- 2. Lu B, Li M, Yin R (2016) Phytochemical content, health benefits, and toxicology of common edible flowers: A review (2000-2015). Critical Reviews in Food Science and Nutrition 56(Sup 1): S130-S148.
- Varzakas T Kiokias S (2016) HPLC Analysis and determination of carotenoid pigments in commercially available plant extracts. Curr Res Nutr Food Sci 4(Special Issue 1): 1-14.
- 4. Wan H, Yu C, Han Y, Guo X, Ahmad S, et al. (2018) Flavonols and carotenoids in yellow petals of rose cultivar (rosa 'sun city'): A possible rich source of bioactive compounds. J Agric Food Chem 66(16): 4171-4181.
- Pavelková P, Krmela A, Schulzová V (2020) Determination of carotenoids in flowers and food supplements by HPLC-DAD. Acta Chimica Slovaca 13(1): 6-12.
- Grajeda IC, Figueroa EMC, Barouh N, Baréa B, Fernandes A, et al. (2016) Isolation and characterization of anthocyanins from *Hibiscus sabdariffa* flowers. Journal of Natural Products 79(7): 1709-1718.
- Wan H, Yu C, Han Y, Guo X, Luo L, et al. (2019) Determination of flavonoids and carotenoids and their contributions to various colors of rose cultivars (*Rosa spp.*), Frontiers in Plant Science 10: 123.
- Kumari P, Raju DVS, Prasad KV, Saha S, Panwar S, et al. (2022) Characterization of anthocyanins and their antioxidant activities in Indian rose varieties (*Rosa×Hybrida*) using HPLC. Antioxidants 11(10): 2032.
- Cendrowski A, Kraśniewska K, Przybył JL, Zielińska A, Kalisz S (2020) Antibacterial and antioxidant activity of extracts from rose fruits (*Rosa rugosa*). Molecules 25(6): 1365.
- 10. Cao X, Xiong X, Xu Z, Zeng Q, He S, et al. (2020) Comparison of phenolic substances and antioxidant activities in different varieties of *chrysanthemum* flower under simulated tea making conditions. Journal of Food Measurement and Characterization 14: 1443-1450.
- 11. Ryu J, Nam B, Kim BR, Kim SH, Jo YD, et al. (2019) Comparative analysis of phytochemical composition of gamma-irradiated mutant cultivars of *chrysanthemum morifolium*. Molecules 24(16): 3003.
- 12. Krzymińska A, Frąszczak B, Gąsecka M, Magdziak Z, Kleiber T (2022) The content of phenolic compounds and organic acids in two *Tagetes patula* cultivars flowers and their dependence on light colour and substrate. Molecules 27(2): 527.
- 13. Yang PF, Yang YN, Feng ZM, Jiang JS, Zhang PC (2019) Six new compounds from the flowers of *chrysanthemum morifolium* and their biological activities. Bioorganic Chemistry 82: 139-144.
- 14. Qiu L, Zhang M, Ju R, Wang Y, Chitrakar B, et al. (2020) Effect of different drying methods on the quality of restructured rose flower (*Rosa rugosa*) chips. Drying Technology 38(12): 1632-1643.

- 15. Villavicencio AL, Heleno SA, Calhelha RC, Santos BC, Barros L, et al. (2018) The influence of electron beam radiation in the nutritional value, chemical composition and bioactivities of edible flowers of *bauhinia variegata l. var. candida alba* Buch-Ham from Brazil. Food Chemistry 241: 163-170.
- 16. Chandler SF, Sanchez C (2012) Genetic modification; the development of transgenic ornamental plant varieties. Plant Biotechnology Journal 10(8): 891-903.
- Azadi P, Bagheri H, Nalousi AM, Nazari F, Chandler SF, et al. (2016) Current status and biotechnological advances in genetic engineering of ornamental plants. Biotechnology Advances 34(6): 1073-1090.
- Jain SM, Spencer MM (2006) Biotechnology and mutagenesis in improving ornamental plants. Floriculture, Ornamental and Plant Biotechnology 1: 589-600.
- 19. Broertjes C (1966) Mutation breeding of *chrysanthemums*. Euphytica 15: 156-162.
- 20. Paramesh TH, Chowdhury S (2005) Impact of explants and gamma irradiation dosage on *in vitro* mutagenesis in carnation (*Dianthus caryophyllus L*). Journal of Applied Horticulture 7(1): 43-45.
- 21. Roychowdhury R, Tah J (2011) Assessment of chemical mutagenic effects in mutation breeding programme for  $M_1$  generation of carnation (*Dianthus caryophyllus*). Research in Plant Biology 1(4): 23-32.
- 22. Datta SK, Chakrabarty D (2005) Classical mutation breeding and molecular methods for genetic improvement of ornamentals. Role of classical mutation breeding in crop improvement. Daya Publishing House, Delhi, India, pp. 260-303.
- 23. Latado R, Hernán AA, Tulmann NA (2004) In vitro mutation of chrysanthemum (Dendranthema grandiflora Tzvelev) with Ethyl Methane Sulphonate (EMS) in immature floral pedicels. Plant Cell, Tissue and Organ Culture 77(1): 103-106.
- 24. Senapati SK, Rout GR (2008) *In vitro* mutagenesis of rose with Ethyl Methane Sulphonate (EMS) and early selection using RAPD Markers. Advances in Horticultural Science 22(3): 218-222.
- 25. Baig MMQ, Hafiz IA, Abbasi NA, Yaseen M, Akram Z, et al. (2012) Reduced-stature rosa species through *in vitro* mutagenesis. Canadian Journal of Plant Science 92(6): 1049-1055.
- 26. Bala M, Pal SK (2013) In vitro mutagenesis of rose (Rosa hybrida L.) explants using gamma-radiation to induce novel flower colour mutations. The Journal of Horticultural Science and Biotechnology 88(4): 462-468.
- 27. Giovannini A, Scariot V, Caser M, Buttafava A, Mansuino A, et al. (2015) Mutation breeding using gamma rays to increase seed germination in *Rosa hybrida*. Acta Horticulturae 1087(1087): 373-378.
- 28. Singh VN, Banerji BK, Dwivedi AK, Verma AK (2009) Effect of gamma irradiation on African marigold (*Tagetes Erecta L.*) Cv. pusa narangi gainda. Journal of Horticultural Sciences 4(1): 36-40.
- 29. Ghosh S, Ganga M, Soorianathasundaram K (2019) Studies on mutagenic effectiveness and efficiency of gamma rays and ethyl methane sulphonate in jasmine. Current Journal of Applied Science and Technology 38(6): 1-8.
- 30. Kumar M, Chaudhary V, Sirohi U, Sharma VR, Naresh RK (2019) Application of molecular markers and their utility in genetic studies of floricultural crops: A review. International Journal of Agriculture Environment and Biotechnology 12(3): 229-247.
- 31. Baliyan D, Sirohi A, Kumar M, Kumar V, Malik S, et al. (2014) Comparative genetic diversity analysis in *chrysanthemum*: A pilot study based on morpho-agronomic traits and ISSR markers. Scientia Horticulturae 167: 164-168.
- 32. Kobeissi B, Saidi A, Kobeissi A, Shafie M (2019) Applicability of SCoT and SSR molecular markers for genetic diversity analysis in *chrysanthemum morifolium* genotypes. Proceedings of the National Academy of Sciences, India Section B: Biological Sciences 89: 1067-1077.

- 33. Chong X, Su J, Wang F, Wang H, Song A, et al. (2019) Identification of favorable SNP alleles and candidate genes responsible for inflorescencerelated traits via GWAS in *chrysanthemum*. Plant Molecul 99: 407-420.
- 34. Agarwal A, GuptaV, Haq SU, Jatav PK, Kothari SL, et al. (2019) Assessment of genetic diversity in 29 rose germplasms using SCoT marker. Journal of King Saud University-Science 31(4): 780-788.
- 35. Senapati SK, Aparajita S, Rout GR (2012) An assessment of genetic fidelity of *in vitro* grown plantlets of rose (*Rosa hybrida*) through molecular markers. African Journal of Biotechnology 11(100): 16532-16538.
- 36. Gaurav AK, Namita, Raju DVS, Ramkumar MK, Singh MK, et al. (2021) Genetic diversity analysis of wild and cultivated Rosa species of India using microsatellite markers and their comparison with morphology based diversity. Journal of Plant Biochemistry and Biotechnology 31: 61-10.
- 37. Yan M, Byrne DH, Klein PE, Yang J, Dong Q, et al. (2018) Genotypingby-sequencing application on diploid rose and a resulting high-density SNP-based consensus map. Horticulture Research 5: 17.
- 38. Ovadis M, Vainstein A, Reuven M, Ben YY, Scovel G (2001) Marker assisted selection for resistance to *Fusarium oxysporum* in the greenhouse carnation. XX International EUCARPIA Symposium, Section Ornamentals, Strategies for New Ornamentals-Part I, Melle, Belgium, pp. 151-156.
- 39. Kimura T, Yagi M, Nishitani C, Onozaki T, Ban Y, et al. (2009) Development of SSR markers in carnation (*dianthus caryophyllus*). Journal of the Japanese Society for Horticultural Science 78(1): 115-123.
- 40. Kumar S, Kumari R, Baheti T, Thakur M, Ghani M (2016) Plant regeneration from axillary bud, callus and somatic embryo in carnation (*Dianthus caryophyllus*) and assessment of genetic fidelity using RAPD-PCR analysis. Indian J Agric Sci 86(11): 1482-1488.
- 41. He YH, Ning GG, Sun YL, Qi YC, Bao MZ (2009) Identification of a SCAR marker linked to a recessive male sterile gene (Tems) and its application in breeding of marigold (*Tagetes erecta*). Plant Breeding 128(1): 92-96.
- 42. Yin ZF, Zhao B, Bi WL, Chen L, Wang QC (2013) Direct shoot regeneration from basal leaf segments of lilium and assessment of genetic stability in regenerants by ISSR and AFLP markers. *In vitro* Cellular & Developmental Biology-Plant 49: 333-342.
- 43. Ramzan F, Kim HT, Younis A, Ramzan Y, Lim KB (2021) Genetic assessment of the effects of self-fertilization in a *Lilium L*. hybrid using molecular cytogenetic methods (FISH and ISSR). Saudi Journal of Biological Sciences 28(3): 1770-1778.
- 44. Mohan Jain S (2002) Feeding the world with induced mutations and biotechnology. pp. 1-14.
- 45. Dobres MS (2011) Prospects for the commercialization of transgenic ornamentals. Transgenic Horticultural Crops, pp. 305-316.
- 46. Rout GR, Mohapatra A, Jain SM (2006) Tissue culture of ornamental pot plant: A critical review on present scenario and future prospects. Biotechnology Advances 24(6): 531-560.
- 47. Passeri V, Koes R, Quattrocchio FM (2016) New challenges for the design of high value plant products: Stabilization of anthocyanins in plant vacuoles. Frontiers in Plant Science 7: 153.
- 48. Tsuda S, Fukui Y, Nakamura N, Katsumoto Y, Yonekura SK, et al. (2004). Flower color modification of *petunia hybrida* commercial varieties by metabolic engineering. Plant Biotechnology 21(5): 377-386.
- 49. Noda N, Yoshioka S, Kishimoto S, Nakayama M, Douzono M, et al. (2017) Generation of blue *chrysanthemums* by anthocyanin B-ring hydroxylation and glucosylation and its coloration mechanism. Science Advances 3(7): e1602785.
- 50. Fukui Y, Tanaka Y, Kusumi T, Iwashita T, Nomoto K (2003) A rationale for the shift in colour towards blue in transgenic carnation flowers expressing the flavonoid 3', 5'-hydroxylase gene. Phytochemistry 63(1):

15-23.

- 51. Qi Y, Lou Q, Quan Y, Liu Y, Wang Y (2013) Flower-specific expression of the phalaenopsis flavonoid 3', 5'-hydoxylase modifies flower color pigmentation in *petunia* and *lilium*. Plant Cell, Tissue and Organ Culture (PCTOC) 115: 263-273.
- Tanaka Y, Brugliera F, Chandler S (2009) Recent progress of flower colour modification by biotechnology. International Journal of Molecular Sciences 10(12): 5350-5369.
- 53. Noman A, Aqeel M, Deng J, Khalid N, Sanaullah T, et al. (2017) Biotechnological advancements for improving floral attributes in ornamental plants. Frontiers In Plant Science 8: 530.
- 54. Lavy M, Zuker A, Lewinsohn E, Larkov O, Ravid U, et al. (2002) Linalool and linalool oxide production in transgenic carnation flowers expressing the *Clarkia breweri* linalool synthase gene. Molecular Breeding 9: 103-111.
- 55. Zvi MMB, Shklarman E, Masci T, Kalev H, Debener T, et al. (2012) PAP1 transcription factor enhances production of phenylpropanoid and terpenoid scent compounds in rose flowers. New Phytologist 195(2): 335-345.
- 56. Brugliera F, Kalc-WG, Hyland C, Webb L, Herbert S, et al. (2000) Improvement of fusarium wilt tolerance in carnations expressing chitinase. Int Plant Mol Biol Rep 18: 522-529.
- 57. Kim YS, Lim S, Yoda H, Choi YE, Sano H (2011) Simultaneous activation of salicylate production and fungal resistance in transgenic *chrysanthemum* producing caffeine. Plant Signaling & Behavior 6(3): 409-412.
- Kumari P, Bhargava B (2021) Phytochemicals from edible flowers: Opening a new arena for healthy lifestyle. Journal of Functional Foods 78: 104375.
- 59. Hong B, Tong Z, Ma N, Li J, Kasuga M, et al. (2006) Heterologous expression of the ATDREB1A gene in *chrysanthemum* increases drought and salt stress tolerance. Science in China Series C: Life Sciences 49: 436-445.
- 60. Chen JR, Lü JJ, Liu R, Xiong XY, Wang TX, et al. (2010) DREB1C from *Medicago truncatula* enhances freezing tolerance in transgenic M. truncatula and China rose (*rosa chinensis jacq.*). Plant Growth Regulation 60: 199-211.
- Matas AJ, Gapper NE, Chung MY, Giovannoni JJ, Rose JK (2009) Biology and genetic engineering of fruit maturation for enhanced quality and shelf-life. Current Opinion in Biotechnology 20(2): 197-203.
- 62. Chandler S (2007) Practical lessons in the commercialization of genetically modified plants-long vase-life carnation. XXVII International Horticultural Congress-IHC2006: International Symposium on Plant Biotechnology: From Bench to Commercialization, Seoul, Korea, pp. 764: 71-82.
- 63. Kaisoon O, Konczak I, Siriamornpun S (2012) Potential health enhancing properties of edible flowers from Thailand. Food Research International 46(2): 563-571.
- 64. Kumari S, Kanth BK, Ahn JY, Kim JH, Lee GJ (2021) Genome-wide transcriptomic identification and functional insight of lily WRKY genes responding to *botrytis* fungal disease. Plants 10(4): 776.
- 65. Ksouda G, Sellimi S, Merlier F, Falcimaigne CA, Thomasset B, et al. (2019) Composition, antibacterial and antioxidant activities of *Pimpinella saxifraga* essential oil and application to cheese preservation as coating additive. Food Chemistry 288: 47-56.
- 66. Kalagatur NK, Mudili V, Kamasani JR, Siddaiah C (2018) Discrete and combined effects of Ylang-Ylang (*Cananga odorata*) essential oil and gamma irradiation on growth and mycotoxins production by *Fusarium graminearum* in maize. Food Control 94: 276-283.
- Malik S (2019) Essential oil research: Trends in biosynthesis, analytics, industrial applications and biotechnological production. Springer, Switzerland.

- 68. Panwar S, Singh KP, Janakiram T, Bhardwaj C, Banyal N, et al. (2017) Molecular characterization of African marigold (*Tagetes erecta*) genotypes using RAPD markers. Indian Journal of Agricultural Sciences 87(5): 663-668.
- 69. Rop O, Mlcek J, Jurikova T, Neugebauerova J, Vabkova J (2012) Edible flowers-a new promising source of mineral elements in human nutrition. Molecules 17(6): 6672-6683.