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Application of epigenetics for allergen-free peanut production: a comprehensive review

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Abstract

Epigenetic methods provide a promising avenue for creating allergen-free peanuts, helping to address the increasing incidence of peanut allergies and related health issues globally. This paper thoroughly explores epigenetic approaches for producing allergen-free peanuts and offers new possibilities for effectively managing peanut allergies. By specifically targeting genes like Arah1, Arah2, and Arah3, the major allergens responsible for allergic reactions can be reduced. Innovative technologies such as CRISPR/dCas9 and RNA interference enable precise regulation of gene activity, facilitating allergen-free peanut production without altering the underlying DNA sequence. The advantages of allergen-free peanuts extend beyond the health benefits, as they have the potential to enhance food safety and open up entirely new markets for peanut-based products. These peanuts could be safely reintroduced into public spaces, like schools and daycare centers, increasing consumer trust and fulfilling the growing demand for safe, nutritious alternatives. However, challenges persist, such as ensuring stable epigenetic changes, preserving other essential peanut traits, and understanding their long-term effects on human health and the environment. Ongoing research should focus on refining these techniques to address these challenges, promote the successful integration of allergen-free peanuts into agricultural systems, and improve public health while transforming the food industry in a sustainable manner.

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Introduction

Peanut allergies are among the most common and serious food allergies, particularly in Western countries, affecting approximately 1%–2% of the population^[1]. These allergies can trigger life-threatening reactions, such as anaphylaxis, which significantly affect individuals and their families due to dietary restrictions and social limitations. The main concern for individuals with peanut allergies is the fear of accidental peanut consumption. The prevalence of peanut allergies has been rising, particularly among children, prompting further investigation into environmental, immunological, and genetic factors that contribute to allergic reactions^[2,3]. In regions such as North America and the United Kingdom, the prevalence of peanut allergies among school-aged children has surpassed 1%, which has raised public health concerns and calls for more research^[4].

The avoidance of peanuts is essential for those affected, as it significantly influences their diet, social activities, and the food industry^[5,6]. The primary proteins in peanuts responsible for triggering allergic reactions are Arah1, Arah2, and Arah3^[7,8]. These allergens provoke immune responses that lead to symptoms like hives, swelling, respiratory distress, and even anaphylaxis. Arah2, in particular, is considered one of the most potent allergens^[9]. Reducing peanut allergenicity presents a challenge, as modifying or removing these proteins without affecting key peanut traits such as nutritional value or taste is complex. Enzymatic hydrolysis has been identified as a potential method for reducing allergenicity; however, chemical modifications may also alter the properties of peanuts, potentially making them more allergenic^[10,11].

Recent advancements in the development of allergen-free peanuts have been achieved through both traditional breeding and cutting-edge gene-editing techniques^[12]. Traditional breeding

efforts have focused on identifying natural peanut varieties with lower allergenic protein content and crossbreeding them to produce cultivars with reduced allergenicity^[11]. The terms 'allergenfree' and 'allergen-reduced' are important in this context. 'Allergenfree' refers to the complete elimination of allergenic proteins, while 'allergen-reduced' means that the quantity or activity of these proteins has been decreased, though they may still be present at lower levels^[11,13].

Epigenetic technologies offer promising strategies for controlling the expression of specific allergenic proteins, such as Arah1, Arah2, and Arah3^[14]. Using tools like CRISPR-based systems, researchers can selectively downregulate the genes responsible for producing these allergens without altering the overall genetic makeup of the peanut^[15,16]. This technique involves modifying DNA methylation or histone modifications at specific loci responsible for allergen expression^[17,18]. Such a targeted approach could reduce peanut allergenicity, leading to peanuts that are safer for individuals with peanut allergies. Practical examples include using epigenetic tools, such as CRISPR-dCas9, to regulate allergen expression in peanuts.

In addition to Arah1, Arah2, and Arah3, Arah6 and Arah8 are also significant contributors to peanut allergies^[19–21]. Arah6 is a 2S albumin, while Arah8 is a profilin, both of which are known allergens, particularly in individuals with latex-fruit syndrome, where there is cross-reactivity between latex and peanut proteins^[22]. Epigenetic tools like CRISPR-dCas9 could be used to target and regulate the expression of these allergens. By modifying the expression of Arah6 and Arah8, it is possible to create peanuts with a reduced allergenic profile, offering safer options for individuals who are sensitive to multiple allergens^[19,23].

Research on the epigenetic regulation of Arah6 and Arah8 is still in its early stages, but initial results are promising. Modifying the expression of these allergens without altering the rest of the peanut genome could lead to peanuts that are safer for individuals with specific sensitivities. Epigenetic silencing of Arah6 and Arah8 could help develop allergen-reduced peanuts that retain vital agronomic traits, such as yield and nutritional value while reducing the risk of allergic reactions^[24].

Despite these advances, challenges persist in developing allergen-reduced peanuts. Limited genetic variability in peanuts and the complexity of allergenic protein profiles remain significant obstacles^[25,26]. However, breeding programs have successfully produced peanut lines with reduced allergen levels, offering safer options for individuals with mild peanut sensitivities. Gene-editing technologies, particularly CRISPR-Cas9, have revolutionized the development of allergen-free peanuts by enabling precise modifications to the genes responsible for allergenic proteins^[15]. Studies have demonstrated the successful silencing of genes such as Arah2 and Arah6, leading to a significant reduction in allergen levels^[27]. This approach accelerates the development process compared to traditional breeding techniques while maintaining specificity and minimizing unintended effects on other peanut traits like yield and flavor.

Furthermore, advancements in molecular biology have enabled the combination of multiple allergen-reducing edits, potentially creating peanuts that are almost completely free of major allergenic proteins^[10]. Although regulatory and public acceptance challenges remain, gene-edited peanuts have shown promise in preclinical studies, with allergenicity tests indicating their potential to be safer for consumption^[28]. These innovations, along with ongoing traditional breeding efforts, offer hope for the widespread adoption of allergen-free peanuts, contributing to safer food systems and addressing the public health burden of peanut allergies.

While significant progress has been made in food science, the management of peanut allergies still relies primarily on avoidance and emergency treatment with epinephrine^[29]. A review of peanut allergy management notes that while avoidance and the use of an epinephrine auto-injector are standard practices, the risk of fatal anaphylaxis or harm from skin contact or inhalation is relatively low^[30]. Research into immunotherapy and desensitization techniques is still in progress, but a definitive solution has not yet been found. The main challenge remains the difficulty of removing or altering allergenic proteins like Arah1, Arah2, and Arah3 without compromising the peanut's nutritional quality or consumer acceptance. Thus, this review aims to explore epigenetic approaches for producing allergen-free peanuts, offering new possibilities for managing peanut allergies and enhancing food safety.

Overview of epigenetics

Epigenetics involves the examination of inheritable changes in gene expression that do not involve modifications to the DNA sequence, driven by mechanisms like DNA methylation, histone alterations, and non-coding RNAs^[31]. In contrast to genetic mutations, epigenetic changes are reversible and can be shaped by environmental factors. Key processes in epigenetics include DNA methylation (which silences gene expression), histone modifications (which modify chromatin structure), and non-coding RNAs (which regulate gene activity after transcription)[32]. Epigenetic mechanisms offer flexible regulation of gene expression, which is essential for processes like development, cellular differentiation, and responses to environmental changes. A review study on precision epigenetic editing: Technological advances, enduring challenges, and therapeutic applications (human studies) states that the epigenome is a complex framework through which gene expression is precisely and flexibly modulated to incorporate heritable

memory and responses to environmental stimuli^[33]. A review of the fundamentals of epigenetics suggests that environmental factors can induce lasting phenotypic changes by causing heritable epigenetic alterations. These modifications influence nucleosomal organization and control whether genes are silenced or activated^[34,35]. Likewise, epigenetics offers the potential to improve crop traits such as stress resistance, disease resilience, and yield by adjusting epigenetic markers, all without altering the plant's DNA. This provides a versatile tool for adapting crops to environmental challenges. Epigenetic editing tools, such as CRISPR/dCas9 and RNA interference, are transforming plant disease treatment and agricultural biotechnology by allowing precise control of gene expression without altering the underlying DNA sequence^[36]. In plant disease management, these technologies can deactivate genes that increase susceptibility to pathogens or activate those that improve disease resistance, providing a targeted and sustainable method for reducing crop losses[37,38]. In agricultural biotechnology, epigenetic editing enhances traits such as stress tolerance, yield, and nutritional guality while also reducing allergenic proteins in crops like peanuts, making them safer for consumption^[39]. This precise regulation of gene activity minimizes unintended effects, offering innovative solutions for managing plant diseases, improving food safety, and promoting agricultural sustainability.

The dendrogram (Fig. 1) illustrates the hierarchical relationships between key epigenetic concepts and their agricultural applications. It emphasizes the connections between genetic regulation, plant disease management, and sustainable farming practices.

Epigenetic editing tools for allergen-free peanut production

Epigenetic editing tools are molecular technologies that precisely modify epigenetic marks, such as DNA methylation and histone modifications, to regulate gene expression without changing the DNA sequence^[40,41].

Epigenetic editing technologies offer innovative possibilities for creating allergen-free peanuts by accurately regulating gene expression without modifying the DNA sequence itself^[42]. A book chapter on CRISPR-Cas technology emphasizes that this system offers several advantages over earlier gene-targeting methods like Transcription Activator-Like Effector Nucleases (TALEN) and Zinc Finger Nucleases (ZFN). These advantages include simpler design, precise genome editing in living cells, higher efficiency, and lower costs^[43]. A promising approach is CRISPR/dCas9-based epigenetic editing, which, unlike conventional CRISPR, does not cut DNA but instead silences specific allergen genes such as Arah1, Arah2, and Arah3 through epigenetic changes, thereby reducing allergen production^[44].

RNA interference technology effectively reduces the expression of allergenic proteins by targeting the mRNA of specific allergen genes, thereby decreasing allergen levels in peanuts. This approach has proven successful in other crops, indicating its potential for application in peanuts^[45,46].

Additionally, emerging epigenetic tools, such as small molecules that modify histones or DNA methylation patterns, are being explored to reduce allergens^[47]. These methods enhance the toolkit for developing hypoallergenic peanut varieties and represent cutting-edge strategies for addressing peanut allergies through epigenetic modification^[48]. Epigenetic editing technologies, such as CRISPR/dCas9, RNA interference, and DNA methyltransferases, allow for precise and heritable suppression of allergenic genes without modifying the DNA sequence, presenting a promising approach for developing allergen-free crops (Table 1).

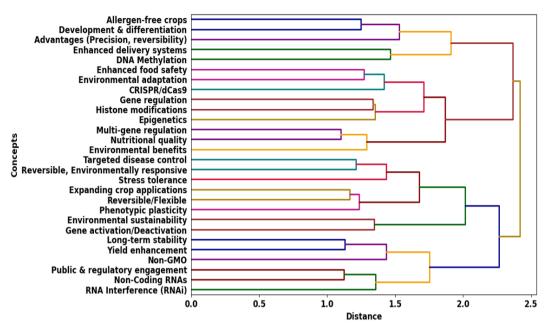


Fig. 1 Hierarchical clustering of epigenetic concepts: unveiling pathways to agricultural innovation.

Table 1. Epigenetic editing tools that could be applied for allergen-free peanut production.

No.	Epigenetic editing tools	Description summary	Ref.
1	CRISPR/dCas9 (Dead Cas9)	Used for targeted gene repression by modifying gene expression without cutting DNA.	[49]
2	Transcription activator	Engineered to precisely target DNA sequences for the epigenetic regulation of gene expression.	[50,51]
3	Zinc finger nucleases	Engineered proteins that target and bind specific DNA sequences to induce epigenetic changes.	[52]
4	RNA interference	Utilizes small RNAs to silence allergenic gene expression at the post-transcriptional level.	[53]
5	DNA Methyltransferase	Used to target and methylate specific gene regions, silencing allergenic genes.	[54]
6	Histone deacetylase	Employed to alter histone acetylation, repressing allergenic gene expression by tightening chromatin.	[55]
7	CRISPR/Cas13	Used to degrade allergenic transcripts in peanuts.	[56]
8	Non-coding RNA-based tools	Regulate the expression of allergenic genes through epigenetic mechanisms.	[57]

Non-transgenic approaches and regulatory considerations

Peanut food allergy is one of the most severe and common food allergies, affecting millions worldwide^[1]. It can trigger life-threatening reactions such as anaphylaxis, characterized by difficulty breathing, a rapid drop in blood pressure, and, in severe cases, death^[58]. Even small amounts of peanut proteins can provoke allergic reactions in sensitive individuals, making this condition a significant public health concern^[59,60]. Peanut allergy often starts in early childhood and tends to persist throughout life, significantly impacting dietary habits and quality of life^[60].

To address peanut allergies, researchers have explored various methods to create allergen-free peanuts. One promising approach involves genetic engineering to silence or modify the genes responsible for producing allergenic proteins^[61,62]. RNA interference (RNAi) technology has been used to suppress the expression of Arah2 and Arah3 proteins, significantly reducing allergenicity. Additionally, breeding techniques, such as traditional crossbreeding or CRISPR-Cas9 genome editing, are being employed to produce hypoallergenic peanut varieties^[63]. Another method involves post-harvest processing, such as enzymatic treatments, to break down allergenic proteins^[64]. These approaches hold great potential for providing safer peanuts while maintaining their nutritional and sensory properties.

Targeted epigenome editing is an advanced technique that enables the precise modification of epigenetic marks, such as DNA methylation or histone modifications, to regulate gene expression without altering the underlying DNA sequence^[40,41]. For example, the review article on epigenome editing technologies for discovery and medicine reports that epigenome editing exploits the heritable and reversible mechanisms of epigenetics to alter gene expression without introducing DNA breaks, inducing DNA damage, or relying on DNA repair pathways^[65]. The approach utilizes tools like CRISPR/dCas9, which pairs a deactivated Cas9 protein with specific epigenetic effector proteins to target specific genomic regions^[66]. Researchers can regulate gene activity by adding or removing specific epigenetic marks, offering a flexible and reversible approach to controlling plant traits. This technique is particularly useful for managing traits governed by complex gene networks, such as improving stress tolerance, increasing yield, or enhancing resistance to diseases^[36].

Applications of targeted epigenome editing in crops have shown immense potential. For example, in rice, researchers have modified epigenetic marks to improve drought and salt tolerance by targeting genes related to stress response pathways^[67]. In tomatoes, epigenome editing has been used to extend shelf life by altering the expression of ripening-related genes^[68,69]. Similarly, in maize, researchers have enhanced yield and biomass production by regulating key genes involved in growth and metabolism^[70,71]. This technique offers an alternative to traditional genetic modification, addressing public concerns about permanent DNA changes while expanding the toolkit for sustainable agriculture. As research advances, epigenome editing could revolutionize crop improvement by offering precise, efficient, and environmentally friendly solutions^[72].

The author of one study suggests that targeted epigenome editing could offer a novel and precise approach to developing allergenfree peanuts by regulating the expression of allergenic genes without altering the DNA sequence^[42]. Tools such as CRISPR/dCas9 combined with epigenetic modifiers can be utilized to suppress the expression of allergen-encoding genes like Arah1, Arah2, and Arah3. This can be achieved by introducing repressive epigenetic marks, such as DNA methylation or histone deacetylation, at the promoters of these genes, thereby significantly reducing or completely silencing their transcription^[6]. This would decrease the production of allergenic proteins, potentially creating peanuts that are hypoallergenic while maintaining their nutritional value and overall quality.

However, there are significant challenges to implementing epigenome editing for allergen-free peanut development (Fig. 2). A major issue is ensuring the long-term stability of the epigenetic modifications across multiple generations, as some marks may be reversible or lost over time^[73]. Another challenge is achieving tissue-specific and precise targeting, as some allergenic proteins may have functional roles in peanut growth and development^[39]. Additionally, off-target effects, where the editing machinery inadvertently affects non-target genes, must be minimized to prevent unintended consequences^[74]. Regulatory and public acceptance of epigenetically modified crops also remains a significant hurdle, as such technologies are still relatively new in agriculture^[75].

Looking ahead, future research should focus on optimizing the delivery systems for epigenome-editing tools, such as nanoparticles or plant transformation methods, to ensure efficient and precise gene targeting in peanuts^[76]. Progress in understanding the epigenetic regulation of allergenic genes will be vital for increasing the consistency and reliability of these modifications. Additionally, combining epigenome editing with other breeding methods, such as CRISPR-Cas9 genome editing or RNA interference, could further improve the efficiency and success of developing allergen-free peanuts^[43]. If these challenges are addressed, targeted epigenome editing could pave the way for safer, allergen-free peanuts and inspire broader applications in addressing food allergies in other crops.

Potential benefits of allergen-free peanuts

Allergen-free peanuts could have a significant public health impact by reducing the prevalence of peanut-related allergic reactions, including life-threatening anaphylaxis. A study on the risks of shared equipment in restaurants for individuals with peanut allergies, specifically in the context of preparing Asian dishes, concludes that an eliciting dose of 300 or 1,000 mg of peanut protein is clinically significant for the peanut-allergic population^[77]. This innovation could greatly reduce the need for strict avoidance measures and emergency treatments for individuals with peanut allergies, improving their overall quality of life and potentially reducing the occurrence of allergies in future generations. The introduction of allergen-free peanuts may also support the growth of peanut-based products and open new market opportunities. A book chapter on the economic and academic significance of peanuts highlights the

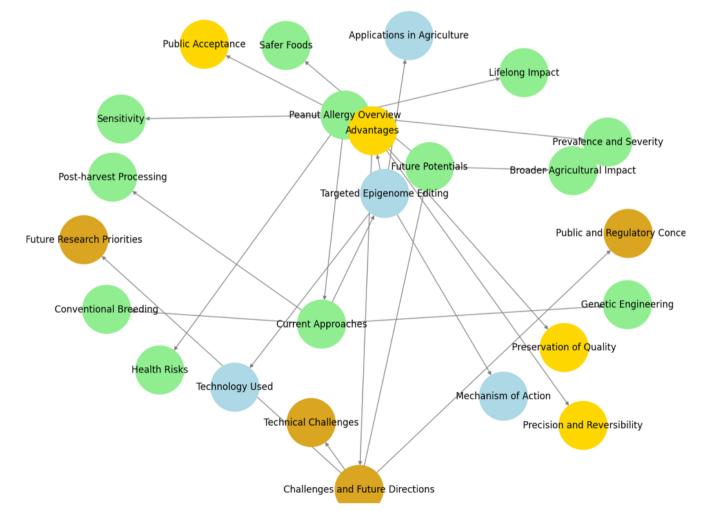


Fig. 2 Exploring approaches for peanut allergy and epigenome editing.

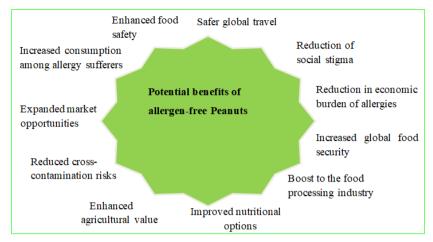


Fig. 3 Health and economic benefits of allergen-free peanuts.

strong global demand for peanuts and peanut-based products, particularly in the confectionery sector^[6]. Reducing concerns about allergic reactions would enable food manufacturers to include peanuts in a wider variety of products, unlocking new opportunities in the food industry^[78]. Educational institutions, airlines, and other organizations could also safely reintroduce peanuts into their offerings, capitalizing on the peanut's versatility and nutritional value. The American Sociological Review article Another Person's Peril: Peanut Allergy, Risk Perceptions, and Responsible Sociality suggests that the three dimensions of proximity to risk material, social, and situational play a key role in shaping policy discussions and influencing how people perceive fear, trust, rights, obligations, and freedom in shared environments, ultimately guiding their views on 'responsible sociality^{1[79]}. Crucially, allergen-free peanuts would preserve their nutritional integrity, providing a sustainable source of protein, healthy fats, and essential nutrients. Research on the bioactive compounds in peanut skin for chronic non-communicable diseases highlights that, due to its polyphenolic content 60 to 120 times higher than that of the peanut grain peanut skin by-products serve as a sustainable source of plant-based bioactives beneficial to human health^[80]. Hence, by mitigating allergens without compromising the peanut's nutritional profile, the peanuts could help to address the increasing global food demands while promoting environmental sustainability in peanut cultivation. Creating allergen-free peanuts holds great promise for improving public health, increasing market accessibility, and minimizing the risk of severe allergic reactions, all while fostering a more inclusive food industry (Fig. 3).

Challenges and future directions

One of the major challenges in producing allergen-free peanuts through epigenetic approaches is achieving stable and consistent modifications. The book chapter 'Managing Food Allergens' proposes a method for controlling allergen risks by setting thresholds that trigger allergic reactions^[81]. Ensuring consistent changes in allergenic gene expression, like Arah1, Arah2, and Arah3, throughout the peanut plant's lifecycle and across generations is a technical challenge, as variability could compromise efforts to reduce allergens.

Another challenge is comprehending the long-term effects and potential reversibility of epigenetic modifications. The Proceedings of the Nutrition Society on Nutritional Developmental Epigenomics: Immediate and Long-Lasting Effects suggests that epigenetic misprogramming during development is believed to have lasting impacts on offspring health, also potentially affecting subsequent

generations^[82]. Since epigenetic modifications do not change the DNA sequence, they can sometimes be reversed by environmental influences or during the plant's reproductive process. Hence, ensuring the stable inheritance of allergen-free traits without unintended effects is crucial for their agricultural viability^[83,84].

Future research efforts should focus on creating more precise and durable epigenetic editing tools, with advancements in technologies such as CRISPR/dCas9 and RNA interference. It is equally important to evaluate the safety and efficiency of these modifications to ensure their viability. Additionally, investigating how environmental factors influence epigenetic alterations and fostering acceptance among consumers and regulatory bodies will be critical steps in advancing the production of allergen-free peanuts.

Conclusions

In conclusion, epigenetics offer a groundbreaking approach to developing allergen-free peanuts by precisely regulating the expression of allergenic genes such as Arah1, Arah2, and Arah3 without altering the underlying DNA sequence. Advanced tools like CRISPR/dCas9 and RNA interference allow for the effective suppression of allergenic protein production, presenting a non-GMO solution to peanut allergies. This innovative application of epigenetics could enhance food safety by reducing allergic reactions, improving the lives of those with peanut allergies, and potentially preventing allergies in future generations. Additionally, allergen-free peanuts could open new market opportunities by making peanuts acceptable in restricted environments such as schools and public spaces. This development would not only boost the peanut industry but also retain the crop's nutritional value and environmental sustainability. Overall, epigenetic strategies have immense potential to revolutionize food safety, improve public health, and strengthen consumer confidence in hypoallergenic food options.

Author contributions

The authors confirm contribution to the paper as follows: draft manuscript preparation, review: Gelaye Y; study supervision, manuscript revision: Luo H. Both authors reviewed the results and approved the final version of the manuscript.

Data availability

Data sharing is not applicable to this article as no new data were analyzed in this study.

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Conflict of interest

The authors declare that they have no conflict of interest.

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References

- Lieberman JA, Gupta RS, Knibb RC, Haselkorn T, Tilles S, et al. 2021. The global burden of illness of peanut allergy: a comprehensive literature review. *Allergy* 76:1367–84
- Tan THT, Ellis JA, Saffery R, Allen KJ. 2012. The role of genetics and environment in the rise of childhood food allergy. Clinical & Experimental Allergy 42:20–29
- Campbell DE, Boyle RJ, Thornton CA, Prescott SL. 2015. Mechanisms of allergic disease–environmental and genetic determinants for the development of allergy. Clinical & Experimental Allergy 45:844–58
- Loh W, Tang MLK. 2018. The epidemiology of food allergy in the global context. International Journal of Environmental Research and Public Health 15:2043
- Du Toit G, Sayre PH, Roberts G, Sever ML, Lawson K, et al. 2016. Effect of avoidance on peanut allergy after early peanut consumption. The New England Journal of Medicine 374:1435–43
- Variath MT, Janila P. 2017. Economic and academic importance of peanut. In *The peanut genome*. Cham: Springer. pp. 7–26. doi:10.1007/978-3-319-63935-2_2
- Jiang S, Wang S, Sun Y, Zhou Z, Wang G. 2011. Molecular characterization of major allergens Ara h 1, 2, 3 in peanut seed. Plant Cell Reports 30:1135–43
- 8. Koppelman SJ, Knol EF, Vlooswijk RAA, Wensing M, Knulst AC, et al. 2003. Peanut allergen *Ara* h 3: isolation from peanuts and biochemical characterization. *Allergy* 58:1144–51
- 9. Palmer GW, Dibbern DA Jr, Wesley Burks A, Bannon GA, Allan Bock S, et al. 2005. Comparative potency of *Ara* h 1 and *Ara* h 2 in immunochemical and functional assays of allergenicity. *Clinical Immunology* 115:302–12
- Shah F, Shi A, Ashley J, Kronfel C, Wang Q, et al. 2019. Peanut allergy: characteristics and approaches for mitigation. Comprehensive Reviews in Food Science and Food Safety 18:1361–87
- Zhou Y, Wang JS, Yang XJ, Lin DH, Gao YF, et al. 2013. Peanut allergy, allergen composition, and methods of reducing allergenicity: a review. *International Journal of Food Science* 2013:909140
- 12. Harshitha T, Parinitha A, Gawali PP, Somya A, Yannam SK. 2024. Grain Processing and Baking Technology. In *Frontiers in Food Biotechnology*. Singapore: Springer. pp. 91–119 . doi: 10.1007/978-981-97-3261-6_7
- Nunavath RS, Singh MT, Chittipolu A, Yerrakula G. 2024. Food processing technology to produce hypoallergenic food with high quality. In Food Allergies: Processing Technologies for Allergenicity Reduction, ed. Lajnaf R. Boca Raton: CRC Press. 348 pp. doi: 10.1201/9781003433781
- Fu G, Zhong Y, Li C, Li Y, Lin X, et al. 2010. Epigenetic regulation of peanut allergen gene Ara h 3 in developing embryos. Planta 231:1049–60
- Brackett NF, Pomés A, Chapman MD. 2022. New frontiers: precise editing of allergen genes using CRISPR. Frontiers in Allergy 2:821107
- Wang M, Schedel M, Gelfand EW. 2024. Gene editing in allergic diseases: identification of novel pathways and impact of deleting allergen genes. *Journal of Allergy and Clinical Immunology* 154:51–58
- Alaskhar Alhamwe B, Khalaila R, Wolf J, von Bülow V, Harb H, et al. 2018. Histone modifications and their role in epigenetics of atopy and allergic diseases. Allergy, Asthma & Clinical Immunology 14:39

- 18. Bégin P, Nadeau KC. 2014. Epigenetic regulation of asthma and allergic disease. *Allergy, Asthma & Clinical Immunology* 10:27
- Kukkonen AK, Pelkonen AS, Mäkinen-Kiljunen S, Voutilainen H, Mäkelä MJ. 2015. Ara h 2 and Ara 6 are the best predictors of severe peanut allergy: a double-blind placebo-controlled study. *Allergy* 70:1239–45
- 20. Hemmings O, Du Toit G, Radulovic S, Lack G, Santos AF. 2020. Ara h 2 is the dominant peanut allergen despite similarities with Ara h 6. *Journal of Allergy and Clinical Immunology* 146:621–630. e5
- 21. Asarnoj A, Nilsson C, Lidholm J, Glaumann S, Östblom E, et al. 2012. Peanut component *Ara* h 8 sensitization and tolerance to peanut. *Journal of Allergy and Clinical Immunology* 130:468–72
- Čelakovská J, Čermákova E, Andrýs C, Boudkova P, Krejsek J. 2024.
 Sensitization to latex and food allergens in atopic dermatitis patients according to ALEX2 Allergy Xplorer test. *Molecular Immunology* 175:89–102
- 23. Bublin M, Breiteneder H. 2014. Developing therapies for peanut allergy. *International Archives of Allergy and Immunology* 165:179–94
- 24. Viquez O, Dodo H, Konan K. 2005. Genetic modification of peanut as a solution to peanut allergy. In *Food Biotechnology*. 2nd Edition. Boca Raton: CRC Press. doi: 10.1201/9781420027976.ch2.17
- 25. Djeghim H, Bellil I, Benslama O, Lekmine S, Temim E, et al. 2024. Effects of genetic diversity on the allergenicity of peanut (*Arachis* hypogaea) proteins: identification of the hypoallergenic accessions using BALB/c mice model and in silico analysis of Ara h 3 allergen cross-reactivity. *Journal of Proteomics* 306:105264
- Cabanillas B, Jappe U, Novak N. 2018. Allergy to peanut, soybean, and other legumes: recent advances in allergen characterization, stability to processing and IgE cross-reactivity. *Molecular Nutrition & Food Research* 62:1700446
- 27. Chu Y, Faustinelli P, Ramos ML, Hajduch M, Stevenson S, et al. 2008. Reduction of IgE binding and nonpromotion of Aspergillus flavus fungal growth by simultaneously silencing Ara h 2 and Ara h 6 in peanut. Journal of Agricultural and Food Chemistry 56:11225–33
- Özgür M, Uçar A. 2024. Food allergen risks in genetically modified foods: current approaches. In Food Safety: Contaminants and Risk Assessment, eds. Kuddus M, Ashraf SA, Rahman P. 1st Edition. Boca Raton: CRC Press. pp. 42–53. di: 10.1201/9781003333913
- 29. Anagnostou K, Clark A. 2015. The management of peanut allergy. Archives of Disease in Childhood 100:68–72
- Restani P, Ballabio C, Corsini E, Fiocchi A, Isoardi P, et al. 2005. Identification of the basic subunit of Ara h 3 as the major allergen in a group of children allergic to peanuts. *Annals of Allergy, Asthma & Immunology* 94:262–66
- Al Aboud NM, Tupper C, Jialal I. 2018. Genetics, epigenetic mechanism.
 In Study Guide. Treasure Island (FL): StatPearls Publishing. https://europepmc.org/article/nbk/nbk532999
- 32. O'Hagan HM. 2014. Chromatin modifications during repair of environmental exposure-induced DNA damage: a potential mechanism for stable epigenetic alterations. *Environmental and Molecular Mutagenesis* 55:278–91
- Roth GV, Gengaro IR, Qi LS. 2024. Precision epigenetic editing: Technological advances, enduring challenges, and therapeutic applications. Cell Chemical Biology 31:1422–46
- 34. Mazzio EA, Soliman KF. 2012. Basic concepts of epigenetics: impact of environmental signals on gene expression. *Epigenetics* 7:119–30
- Conner JA, Guimaraes LA, Zhang Z, Marasigan K, Chu Y, et al. 2024.
 Multiplexed silencing of 2S albumin genes in peanut. *Plant Biotechnology Journal* 22:2438–40
- Miglani GS, Kaur A, Kaur L. 2020. Plant gene expression control using genome-and epigenome-editing technologies. *Journal of Crop Improve*ment 34:1–63
- 37. van Esse HP, Reuber TL, van der Does D. 2020. Genetic modification to improve disease resistance in crops. *New Phytologist* 225:70–86
- 38. Tirnaz S, Batley J. 2019. DNA methylation: toward crop disease resistance improvement. *Trends in Plant Science* 24:1137–50
- Phogat S, Lankireddy SV, Lekkala S, Anche VC, Sripathi VR, et al. 2024.
 Progress in genetic engineering and genome editing of peanuts: revealing the future of crop improvement. *Physiology and Molecular Biology of Plants* 30:1759–75
- Thakore PI, Black JB, Hilton IB, Gersbach CA. 2016. Editing the epigenome: technologies for programmable transcription and epigenetic modulation. *Nature Methods* 13:127–37

- Seem K, Kaur S, Kumar S, Mohapatra T. 2024. Epigenome editing for targeted DNA (de)methylation: a new perspective in modulating gene expression. Critical Reviews in Biochemistry and Molecular Biology 59:69–98
- 42. Sharma Y, Patil P, Khatal MM, Huchchannanavar S, Rachana MN, et al. 2024. Biological and chemical factors influencing food allergies: a comprehensive review. *Journal of Scientific Research and Reports* 30:787–94
- Mohan C, Satish L, Muthubharathi BC, Selvarajan D, Easterling M, et al. 2022. CRISPR-Cas technology: A genome-editing powerhouse for molecular plant breeding. In *Biotechnological innovations for environ*mental bioremediation, eds. Arora S, Kumar A, Ogita S, Yau YY. Singapore: Springer. pp. 803–79. doi: 10.1007/978-981-16-9001-3_32
- 44. Rajanathadurai J, Perumal E, Sindya J. 2024. Advances in targeting cancer epigenetics using CRISPR-dCas9 technology: A comprehensive review and future prospects. *Functional & Integrative Genomics* 24:164
- 45. Konan KN, Ananga AO, Dodo H. 2015. Reducing peanut allergy risks by means of genetic modification. In *Modification of Seed Composition to Promote Health and Nutrition*. Madison, WI, USA: American Society of Agronomy, Crop Science Society of America, Soil Science Society of America. pp. 289–302. doi: 10.2134/agronmonogr51.c15
- 46. White BL, Shi X, Burk CM, Kulis M, Wesley Burks A, et al. 2014. Strategies to mitigate peanut allergy: production, processing, utilization, and immunotherapy considerations. *Annual Review of Food Science and Technology* 5:155–76
- 47. Potaczek DP, Alashkar Alhamwe B, Miethe S, Garn H. 2021. Epigenetic mechanisms in allergy development and prevention. In *Allergic diseases-from basic mechanisms to comprehensive management and prevention*. Springer, pp. 331–57. doi: 10.1007/164_2021_475
- 48. Mousavi Khaneghah A, Mostashari P. 2024. Decoding food reactions: a detailed exploration of food allergies vs. intolerances and sensitivities. Critical Reviews in Food Science and Nutrition 65(14):2669–713
- Kanafi MM, Tavallaei M. 2022. Overview of advances in CRISPR/dead-Cas9 technology and its applications in human diseases. Gene 830:146518
- 50. Pelletier S. 2016. Genome Editing with Targetable Nucleases. In *Genome Editing*. Cham: Springer. pp. 1-29. doi: 10.1007/978-3-319-34148-4_1
- Parkhi V, Bhattacharya A, Char B. 2021. Multiomics technologies and genetic modification in plants: rationale, opportunities and reality. In Omics Technologies for Sustainable Agriculture and Global Food Security (Vol II), ed. Kumar A, Kumar R, Shukla P, Patel HK. pp. 313–28. doi: 10.1007/978-981-16-2956-3_12
- 52. Gersbach CA, Gaj T, Barbas CF 3rd. 2014. Synthetic zinc finger proteins: the advent of targeted gene regulation and genome modification technologies. *Accounts of Chemical Research* 47:2309–18
- 53. Downward J. 2004. RNA interference. *BMJ* 328:1245–48
- Lee KH, Song Y, O'Sullivan M, Pereira G, Loh R, et al. 2017. The implications of DNA methylation on food allergy. *International Archives of Allergy and Immunology* 173:183–92
- Bhavsar P, Ahmad T, Adcock IM. 2008. The role of histone deacetylases in asthma and allergic diseases. *Journal of Allergy and Clinical Immunol*ogy 121:580–84
- 56. Adeyinka OS, Tabassum B, Koloko BL, Ogungbe IV. 2023. Enhancing the quality of staple food crops through CRISPR/Cas-mediated site-directed mutagenesis. *Planta* 257:78
- 57. Weinberg MS, Morris KV. 2013. Long non-coding RNA targeting and transcriptional de-repression. *Nucleic Acid Therapeutics* 23:9–14
- 58. Reyes AJ, Hosein AS, Ramcharan K, Perot S. 2020. Anaphylaxis and other allergic reactions to food: a global challenge. *BMJ Case Reports* 13:e231425
- 59. Gray CL. 2020. Current controversies and future prospects for peanut allergy prevention, diagnosis and therapies. *Journal of Asthma and Allergy* 13:51–66
- Muthukumar J, Selvasekaran P, Lokanadham M, Chidambaram R. 2020.
 Food and food products associated with food allergy and food intolerance–An overview. Food Research International 138:109780
- 61. Dodo H, Konan K, Viquez O. 2005. A genetic engineering strategy to eliminate peanut allergy. *Current Allergy and Asthma Reports* 5:67–73
- Singh MB, Bhalla PL. 2008. Genetic engineering for removing food allergens from plants. *Trends in Plant Science* 13:257–60
- 63. Abady S, Shimelis H, Janila P, Mashilo J. 2019. Groundnut (*Arachis hypogaea* L.) improvement in sub-Saharan Africa: a review. *Acta Agriculturae Scandinavica*, *Section B Soil & Plant Science* 69:528–45

- 64. Chizoba Ekezie FG, Cheng JH, Sun DW. 2018. Effects of nonthermal food processing technologies on food allergens: a review of recent research advances. *Trends in Food Science & Technology* 74:12–25
- McCutcheon SR, Rohm D, Iglesias N, Gersbach CA. 2024. Epigenome editing technologies for discovery and medicine. *Nature Biotechnology* 42:1199–217
- 66. Tadić V, Josipovi ć G, Zoldoš V, Vojta A . 2019. CRISPR/Cas9-based epigenome editing: an overview of dCas9-based tools with special emphasis on off-target activity. *Methods* 164:109–19
- 67. Li Q, Zhu P, Yu X, Xu J, Liu G. 2024. Physiological and molecular mechanisms of rice tolerance to salt and drought stress: advances and future directions. *International Journal of Molecular Sciences* 25:9404
- 68. Ming Y, Jiang L, Ji D. 2023. Epigenetic regulation in tomato fruit ripening. Frontiers in Plant Science 14:1269090
- 69. Chen S, Han J, Wu S, Guo S, Tang Y, et al. 2024. From non-coding RNAs to histone modification: The epigenetic mechanisms in tomato fruit ripening and quality regulation. *Plant Physiology and Biochemistry* 215:109070
- 70. Li Z, Gao Q, Liu Y, He C, Zhang X, et al. 2011. Overexpression of transcription factor ZmPTF1 improves low phosphate tolerance of maize by regulating carbon metabolism and root growth. *Planta* 233:1129–43
- Long Y, Wang C, Liu C, Li H, Pu A, et al. 2024. Molecular mechanisms controlling grain size and weight and their biotechnological breeding applications in maize and other cereal crops. *Journal of Advanced Research* 62:27–46
- Kakoulidou I, Avramidou EV, Baránek M, Brunel-Muguet S, Farrona S, et al. 2021. Epigenetics for crop improvement in times of global change. *Biology* 10:766
- 73. Trerotola M, Relli V, Simeone P, Alberti S. 2015. Epigenetic inheritance and the missing heritability. *Human Genomics* 9:17
- 74. Movahedi A, Aghaei-Dargiri S, Li H, Zhuge Q, Sun W. 2023. CRISPR variants for gene editing in plants: biosafety risks and future directions. International Journal of Molecular Sciences 24:16241
- 75. Caradus JR. 2023. Processes for regulating genetically modified and gene edited plants. *GM Crops & Food* 14:1–41
- Vats S, Kumawat S, Kumar V, Patil GB, Joshi T, et al. 2019. Genome editing in plants: exploration of technological advancements and challenges. Cells 8:1386
- Remington BC, Blom WM, Bassa B, Koppelman SJ. 2020. Risk of shared equipment in restaurants for consumers with peanut allergy: a simulation for preparing Asian foods: a simulation for preparing Asian foods. Annals of Allergy, Asthma & Immunology 125:543–551.e6
- Sicherer SH, Sampson HA. 2007. Peanut allergy: Emerging concepts and approaches for an apparent epidemic. *Journal of Allergy and Clinical Immunology* 120:491–503
- DeSoucey M, Waggoner MR. 2022. Another person's peril: peanut allergy, risk perceptions, and responsible sociality. American Sociological Review 87:50–79
- 80. Cordeiro-Massironi K, Freitas RAMS, da Silva Martins ICV, de Camargo AC, da Silva Torres EAF. 2024. Bioactive compounds of peanut skin in prevention and adjunctive treatment of chronic non-communicable diseases. Food & Function 15:6304–23
- 81. Walker MJ, Gowland MH. 2017. Food allergy: managing food allergens. In *Analysis of Food Toxins and Toxicants*, eds. Wong YC, Lewis RJ. United Kingdom: John Wiley & Sons. pp.711–42. doi: 10.1002/9781118992685. ch24
- 82. Attig L, Gabory A, Junien C. 2010. Nutritional developmental epigenomics: immediate and long-lasting effects. *The Proceedings of the Nutrition Society* 69:221–31
- Dwivedi S, Puppala N, Maleki S, Ozias-Akins P, Ortiz R. 2016. Peanut improvement for human health. In *Plant Breeding Reviews*, eds. Janick J. vol. 38. Hoboken, New Jersey: Wiley-Blackwell. pp. 143–86. doi: 10.1002/9781118916865.ch04
- Lee NA, Wright GC, Rachaputi RC. 2016. Peanuts: Bioactives and allergens.
 USA: DEStech Publications, Inc. www.destechpub.com/wp-content/uploads/2016/03/Peanuts-Bioactive-Allergens-preview.pdf?x74822



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