

Murine Models of Allergic Asthma: Methodological Insights into Allergen Sensitization and Challenge Protocols

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ABSTRACT

Asthma represents a chronic inflammatory airway disease with a steadily increasing global prevalence in recent decades. Animal models have proven invaluable in elucidating the underlying disease mechanisms and identifying innovative therapeutic approaches. The murine model is extensively used to investigate key characteristics of allergic asthma, including airway inflammation, airway hyperresponsiveness (AHR), and airway remodeling. Classic protocols involving sensitizing and challenging animals with different types of allergens and modes of administration are major factors in inducing asthmatic features in a mouse model. The present review critically analyzes the commonly used sensitization and allergen challenge protocols for inducing acute and chronic inflammation in the airways of mouse models of asthma, emphasizing their potential in advancing therapeutic development for allergic asthma studies.

Key words: Asthma, acute mouse model, chronic mouse model, sensitization, and challenge

INTRODUCTION

Asthma affects approximately 300 million people globally and continues to exhibit a rising trend every year¹. Its intricate nature arises from a complex interplay of genetic and environmental factors². Allergic asthma, the typical phenotype in clinical asthma, is triggered by allergen exposure, manifesting as a chronic inflammatory disorder affecting the airways. Key features of asthma include airway inflammation, eosinophilia, goblet cell hypersecretion, airway hyperresponsiveness (AHR), and airway remodeling³.

The cellular and biochemical processes underlying the development of allergic airways, associated with airway inflammation and remodeling, have been investigated in clinical and animal studies⁴. Studying asthma in humans is ethically challenging, although it is the best approach to understand the pathophysiology of the disease and to investigate drug efficacy for new drug development in allergic asthma. Hence, the utilization of animal models is essential for a comprehensive understanding of the disease, notwithstanding their limitations in replicating the complexity of human asthma.

The mouse model is widely employed to investigate the involvement of various cells and mediators, as well as structural and physiological manifestations of allergic asthma progression. This review focuses on the establishment of allergic asthma, incorporating different types of allergens and administration

methods during sensitization and challenge in an asthmatic mouse model.

ALLERGIC-INDUCED TYPE 2 EOSINOPHILIC ASTHMA

In general, asthma is categorized into type 2 and non-type 2 inflammation based on distinct endotypes (Figure 1). Airway inflammation in type 2 immune response-driven asthma is phenotypically expressed as eosinophilic asthma, while non-type 2 immune response-driven asthma is characterized as neutrophilic asthma and paucigranulocytic asthma⁵.

Eosinophilic asthma is marked by increased eosinophil production and infiltration in the airways in response to an allergen. In type 2 immune response-driven asthma, the increase in T helper 2 (Th2) lymphocytes in the peripheral blood of asthmatic patients during an exacerbation is related to the severity of airway eosinophilia, contributing to the pathophysiological changes that require aggressive treatment⁶. Upon contact with allergens presented by antigen-presenting cells (APCs) in the airway, Th2 cells secrete Th2 cytokines such as interleukin (IL)-4, IL-5, and IL-13, which recruit inflammatory cells (including eosinophils, basophils, and mast cells) and activate B cells to release immunoglobulin E (IgE) (Figure 2)⁷.

IL-13 targets goblet cells, leading to excessive mucus production and goblet cell hyperplasia; it also

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History

- Received: 12-6-2024
- Accepted: 06-4-2025
- Published Online: 30-4-2025

DOI : 10.15419/bmrat.v12i4.973



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Cite this article : Mohd Rosdan Bushra S, Abdullah Nurul A. Murine Models of Allergic Asthma: Methodological Insights into Allergen Sensitization and Challenge Protocols. *Biomed. Res. Ther.* 2025; 12(4):7320-7334.

induces eosinophil infiltration by priming the vessel wall, resulting in AHR⁸. IL-5 participates in the development, activation, and migration of eosinophils from the bone marrow to the airways, initiating airway inflammation⁹. IL-4 initiates IgE isotype class switching in B cells and upregulates the IgE receptor (FcεRI) on the mast cell surface, resulting in the release of histamine and other mediators³. Another hallmark of asthma is the elevated level of serum IgE synthesized by plasma cells activated by IL-4-induced class switching of B cells¹⁰.

MOUSE STRAIN IN THE ALLERGIC ASTHMA MOUSE MODEL

Mouse strains exhibit diverse capabilities in manifesting specific diseases and play a major role in ensuring the successful development of intended phenotypes. The most widely preferred strains include *BALB/c*, *C57BL/6*, and *A/J* mice¹¹.

The *BALB/c* strain has become particularly prominent in asthma studies involving allergen challenge due to its proficiency in activating a robust type 2 immune response¹². This includes the production of Th2 cytokines, allergen-specific IgE, eosinophilic responses, and AHR. Upon allergen exposure, *BALB/c* mice readily produce these Th2 cytokines and develop AHR and airway inflammation, characterized by eosinophilic infiltration—all of which are crucial to the pathogenesis of allergic asthma^{13,14}. Furthermore, their strong tendency to produce IgE antibodies in response to allergens facilitates the sensitization phase of allergic asthma. Upon allergen re-exposure, the crosslinking of IgE to mast cells subsequently triggers degranulation and the release of inflammatory mediators^{15,16}. Moreover, *BALB/c* mice exhibit airway remodeling, demonstrating their capability to express the pathophysiology of the inflammatory process in asthma¹⁷.

In contrast, the *C57BL/6* strain is regarded as a prototypic non-type 2 mouse strain, eliciting Th1 cytokines (interferon-gamma (IFN- γ) and tumor necrosis factor alpha (TNF- α)) in response to allergen challenge¹⁸. Despite limitations in allergic airway development—particularly in IgE expression and AHR to methacholine—this strain is widely employed as a genetically modified animal model for assessing the impact of genetic manipulation on disease progression, including evaluation of allergen sensitization responsiveness and allergic airway inflammation¹⁹. Researchers also utilize other strains,

such as *A/J*, in mouse models of asthma, demonstrating effectiveness in inducing AHR and increasing cytokine production²⁰.

Nevertheless, mouse models of asthma do not perfectly recapitulate the complexity of human asthma, largely due to the heterogeneous nature of the disease with various phenotypes. Modeling the full spectrum of human asthma in a single mouse is challenging, often necessitating a focus on specific mechanisms, such as Th2-mediated inflammation and AHR²¹. Significant differences exist between mouse and human airway anatomy and physiology, including variations in size, structure, and branching patterns that affect allergen delivery and the development of airway inflammation and remodeling²². While the mouse and human immune systems share similarities, genetic variations lead to significant differences, particularly in cytokine profiles, receptor expression, and gene regulation, which influence asthma development and progression²³. Furthermore, artificial allergen sensitization protocols commonly used in mouse models—often involving repeated exposure to high doses of purified allergens—differ from natural human allergen exposure, which is typically more chronic and involves a complex mixture of allergens²⁴.

ALLERGENS USED TO INDUCE ASTHMA

An allergen is any substance recognized as foreign by the immune system, provoking an allergic response. Different types of allergens can induce asthmatic conditions in animal models, with ovalbumin (OVA) being a commonly employed allergen. Whether in acute or chronic models, OVA offers advantages such as affordability, availability, a highly purified antigen, well-defined major histocompatibility complex (MHC) epitopes, and the existence of a recombinant peptide, making it a popular choice²⁵. The allergic reaction induced by OVA produces a rapid, strong, and standardized response. The OVA-sensitized and challenged mouse models have successfully elucidated the effects of inflammatory cell infiltration, Th2 cytokine secretion, eosinophil recruitment, AHR, and airway remodeling²⁶. Additionally, some studies have reported goblet cell hyperplasia, increased mucus production, collagen deposition, and fibrosis²⁷.

While the OVA model has greatly contributed to understanding the mechanisms of allergic asthma, concerns persist regarding its clinical relevance. Challenges in using OVA-induced asthma models include

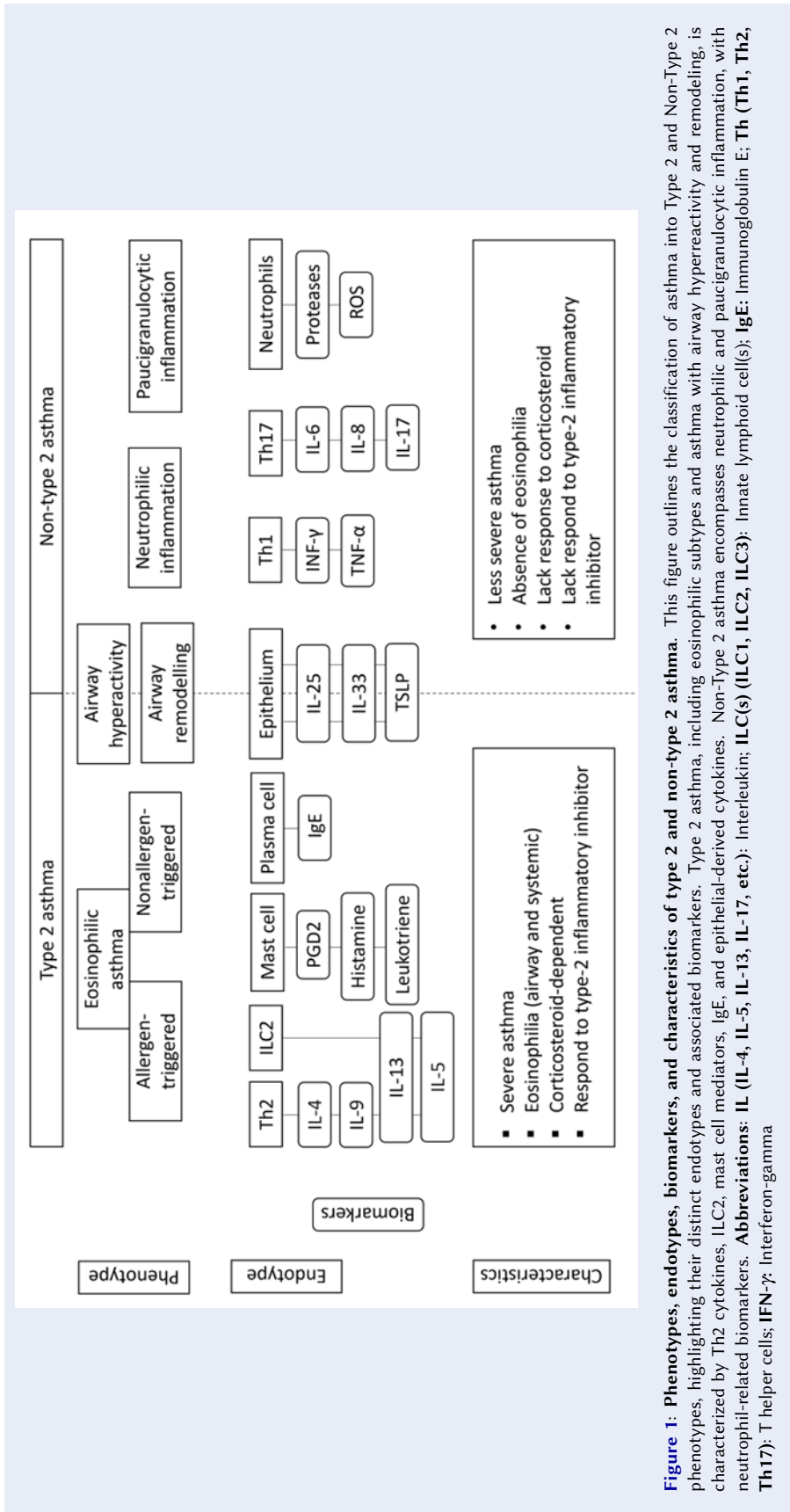


Figure 1: Phenotypes, endotypes, biomarkers, and characteristics of type 2 and non-type 2 asthma. This figure outlines the classification of asthma into Type 2 and Non-Type 2 phenotypes, highlighting their distinct endotypes and associated biomarkers. Type 2 asthma, including eosinophilic subtypes and asthma with airway hyperreactivity and remodeling, is characterized by Th2 cytokines, ILC2, mast cell mediators, IgE, and epithelial-derived cytokines. Non-Type 2 asthma encompasses neutrophilic and paucigranulocytic inflammation, with neutrophil-related biomarkers. **Abbreviations:** IL (IL-4, IL-5, IL-13, IL-17, etc.): Interleukin; ILC(s) (ILC1, ILC2, ILC3): Innate lymphoid cell(s); IgE: Immunoglobulin E; Th (Th1, Th2, Th17): T helper cells; IFN- γ : Interferon-gamma

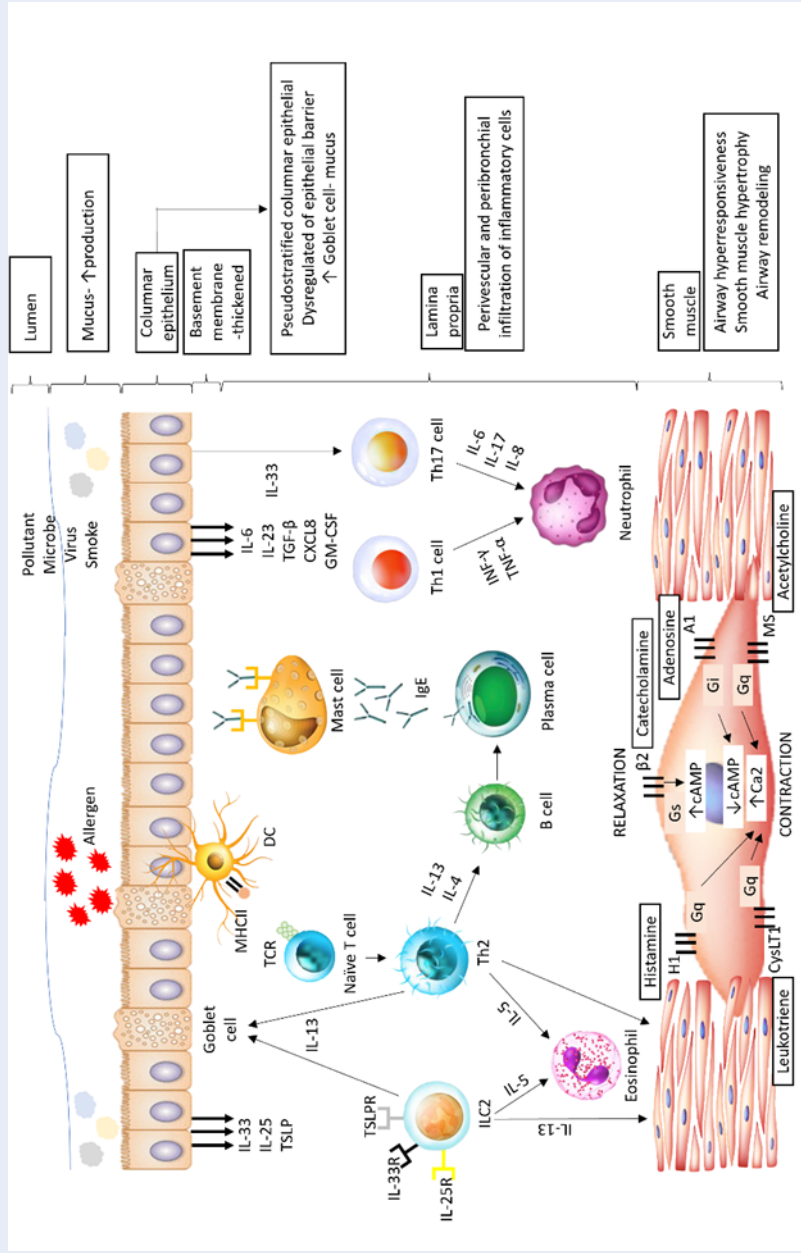


Figure 2: The orchestrated pathways of asthma exacerbation involve inflammatory cell infiltration, cytokine production, AHR, and airway remodeling. This diagram illustrates the immune mechanisms involved in asthma. Allergens and other stimuli trigger the release of cytokines from the epithelium, activating ILC2 and Th2 cells. These cells promote eosinophil recruitment and IgE production by plasma cells, leading to mast cell degranulation and the release of histamine and leukotrienes, causing airway inflammation and smooth muscle contraction. Non-Th2 pathways involving Th1 and Th17 cells and neutrophils also contribute to inflammation and airway remodeling. **Abbreviations:** IFN- γ : Interferon-gamma, IgE: Immunoglobulin E, IL (IL-4, IL-5, IL-13, IL-17, etc.): Interleukin, ILC(s) (ILC1, ILC2, ILC3): Innate lymphoid cell(s), Th (Th1, Th2, Th17): T helper cells, TNF- α : Tumor necrosis factor alpha, and TSLP: Thymic stromal lymphopoietin

the development of OVA tolerance during long-term interventions in chronic models, the discrepancy between the human (airway) and mouse (intraperitoneal) sensitization routes—which may bypass the innate airway immune environment—and the rarity of encountering OVA in human asthma²⁸. Consequently, other models using allergens more closely related to human asthma, such as house dust mite (HDM), have been developed.

Dermatophagoides farinae (American HDM) and *Dermatophagoides pteronyssinus* (European HDM) are common aeroallergens known to cause allergic sensitization²⁹. HDM inhalation triggers pattern recognition receptors (PRRs) on airway epithelial cells, leading to chemokine and cytokine secretion that cause damage to the airway epithelia³⁰. The allergenicity of HDM depends on its allergenic protein load, reflected by the IgE-binding complex pattern measured by the antibody titer.

HDM sensitization and challenge in mouse models have successfully reproduced asthmatic features³¹. Immunotherapy with purified natural *D. pteronyssinus* reduced AHR, eosinophilia, and Th2 cytokines in mice, indicating potential clinical effects³². Derp 2.1 peptide treatment has demonstrated the ability to suppress Th2 and Th17 cell polarization via IL-10-secreting dendritic cells³³. Derp2-FlaB fusion protein, used as a treatment in HDM-sensitized mice, inhibited AHR, eosinophil infiltration, and Derp2-specific IgE, suggesting promise as a vaccine in asthma therapy³⁴.

Additionally, other allergens have also been used for sensitization and challenge in asthmatic mouse models. Acute allergic inflammation induced by papain was observed to stimulate eosinophilia⁹. Intratracheal challenge with *Schizophyllum commune* fungus in an OVA-induced model increased airway neutrophilia and the secretion of IL-17A and IL-17F³⁵. Coal fly dust used to sensitize *BALB/c* mice enhanced neutrophil and other inflammatory cell infiltration, as well as increased cytokine secretion³⁶. Sensitization and challenge in mice using shrimp tropomyosin resulted in eosinophilia, increased IgE secretion, lung inflammation, mucus hypersecretion, goblet cell hyperplasia, collagen deposition, and dense smooth muscle, indicating that shrimp tropomyosin can be employed as an allergen to study asthma pathogenesis³⁷.

ALLERGEN SENSITIZATION IN MOUSE MODELS

Sensitization procedures are essential for inducing asthmatic conditions in animal models. Since

asthma does not naturally develop in mice, sensitization is necessary to introduce the allergen and requires multiple re-exposures to evoke the allergic reaction. The initial exposure to the allergen stimulates T lymphocytes to secrete Th2 cytokines, while B lymphocytes undergo isotype switching, generating allergen-specific IgE³⁸. Subsequent re-exposures lead to the cross-linking of basophils and IgE-bound mast cells, triggering degranulation and the release of inflammatory mediators.

Allergens are commonly used to induce allergic responses in animal models, together with adjuvants to enhance the immunogenicity of the allergen and further support the development of asthmatic animal models³⁹. OVA, HDM, and *Aspergillus* are clinically relevant allergens in humans and are commonly used in allergic asthma mouse models. OVA, a protein allergen mainly found in chicken's egg white, is widely used in the majority of studies on allergic asthma⁴⁰. Various routes of sensitization, including intraperitoneal (i.p.), subcutaneous (s.c.), intranasal (i.n.) injection, and epicutaneous (ec), can be used to induce asthmatic conditions.

In the development of animal allergic asthma models, an adjuvant is administered to enhance the sensitization mechanism of allergens during the sensitization phase. Aluminum hydroxide (alum), frequently used as an adjuvant, induces a strong type 2 immune reaction⁴¹. The aggregate structure of alum continuously releases antigen, promoting phagocytosis and inducing local inflammation, resulting in macrophage activation, MHC class II expression, and antigen presentation⁴². The recruitment of macrophages and dendritic cells was observed in the alum-adjuvant group, with increased eosinophilic infiltration, Th2 cytokines, and IgE levels⁴³.

In contrast, Complete Freund's Adjuvant (CFA) induces Th17 and Th1 cell activation, resulting in neutrophilic infiltration of the lungs⁴⁴. A few studies have reported significant neutrophil infiltration and low eosinophil numbers, indicating that CFA is effective in inducing neutrophilic asthma⁴⁵. The lungs were also dominated by dendritic cells, macrophages, and activated B cells, with increases in the Th1 cytokine IFN γ and the Th17 cytokine IL-17A⁴³. Interestingly, different allergens administered with the same adjuvant produced different effects, where subcutaneous injection of OVA/CFA showed neutrophilic inflammation⁴⁶, whereas HDM/CFA exhibited mixed eosinophilic-neutrophilic inflammation⁴⁷. This difference is possibly due to the distinct nature of the antigens and how they interact with the immune system. When

combined with strong adjuvants like CFA, OVA, a relatively simple protein, may preferentially stimulate a robust Th1 immune response⁴⁸, whereas HDM, a complex mixture of proteins, can activate a broader immune response, engaging both Th2 and Th17 cells⁴⁹.

Meanwhile, lipopolysaccharide (LPS) is widely used to induce mixed eosinophilic and neutrophilic inflammation in asthmatic mouse models⁵⁰. LPS activates toll-like receptor 4 (TLR4) on lung epithelial cells, transducing a pro-inflammatory signaling pathway⁵¹. The concentration of inhaled LPS during sensitization determines the type of inflammation, where low levels of LPS lead to Th2 responses, while high levels induce Th1 responses⁵².

Nevertheless, the use of adjuvants can alter experimental animal behavior by causing distress and interfering with the study of adjuvant-containing drugs, such as allergen-specific immunotherapy for allergy vaccine development⁵³. Hence, adjuvant-free sensitization offers a more realistic model, mirroring chronic asthma manifestation in humans⁵⁴. Adjuvant-free sensitization via subcutaneous injection can induce AHR, airway remodeling, increased IgE secretion, and eosinophil and lymphocyte infiltration⁵⁵. Likewise, the intranasal route can induce allergic inflammation associated with Th2 cytokine secretion, increased inflammatory cell infiltration, and mucus hypersecretion⁵³.

Therefore, the route of sensitization, as well as the types of adjuvants and allergens used, play pivotal roles in inducing different phenotypes of asthma inflammation. The presence of various adjuvants in allergen sensitization leads to different inflammatory responses in the asthmatic airway (**Table 1**).

ALLERGEN CHALLENGE IN MOUSE MODELS

The capability of mouse models to induce the asthmatic condition is well-established, and these models are useful for controlling inflammation. The acute allergic airway inflammatory model is predominantly studied due to its ability to successfully establish many asthmatic features. However, this acute model falls short in developing other major features observed in human asthma, such as collagen deposition and chronic airway remodeling. Consequently, the field has shifted toward developing and studying chronic allergic airway inflammation models to address the limitations of the acute model.

Acute allergen challenge model

Because mice do not naturally develop asthma, human intervention is necessary to induce artificial asthmatic conditions in the airways. Asthma is characterized by multiple phenotypes and cannot be entirely replicated by a single model. Hence, specific phenotypes are developed depending on the objectives of the study. **Table 2** provides a summary of different sensitization and challenge protocols in acute asthmatic mouse models.

The development of an asthmatic model in mice depends on several factors, including the protocol of sensitization and challenges, the adjuvants, and the type of allergens. In the acute mouse asthma model, diverse yet coherent protocols were employed. Allergen sensitization via systemic delivery into the circulatory system commonly necessitates multiple re-exposures to establish a favorable allergic model³⁸. Meanwhile, allergen challenge is usually administered via the airways through inhalation (aerosol), intratracheal (i.t.), or intranasal (i.n.) routes. The common acute model protocol involves allergen sensitization lasting for two to three weeks, followed by allergen challenge for several consecutive days, with the endpoint assessed 24 hours after the last challenge.

The acute mouse model develops the common characteristics of clinical asthma. Studies have shown that lung pathology induced by allergens can exhibit changes in the lungs that cause airway inflammation, airway remodeling, and AHR^{66?}. Histological analysis allows examination of inflammatory cell recruitment, mucus production, collagen deposition, and fibrosis in the perivascular and peribronchiolar space⁶⁷. The acute model is also utilized to study the mechanisms of remodeling and oxidative stress associated with the signaling pathway in pulmonary asthma^{68,69}. Additionally, this model has also shown the amelioration of allergic inflammation when treated with various potential suppressors, such as IL-38⁷⁰, anti-IL-25⁷¹, and leukotriene B4 receptor blocker⁷².

While the acute model has successfully investigated some features of the pathophysiology of asthma, it has limitations compared to clinical asthma, which requires persistent airway inflammation to mimic asthmatic individuals. The short period of allergen challenge is one reason for minimal changes in airway remodeling, AHR, and eosinophilia, with these changes subsiding a few weeks after the last challenge. Asthma is associated with chronic disease, so some concerns arise regarding the reliability of acute mouse models in investigating disease progression and potential treatments.

Table 1: The route of allergen and adjuvant sensitization and its effect in asthma development

Allergen	Adjuvant	Strain	Route	Efficacy	Reference
OVA	Alum	BALB/c	i.p.	↑ eosinophils and B cells population ↓ GATA3 and ILC2s in LN ↓ IFN- γ and Th1 cells in lung ↑ IL-5 and IL-4 and Th2 cells in lung and LN	52
	LPS	BALB/c	i.p.	↓ eosinophils percentage ↑ neutrophils population in BALF ↑ T-bet and ILC1s in lungs ↑ ROR γ t and ILC3s in LN ↑ Th17 cells in lungs and LN	
OVA	CFA	C57BL/6	i.p.	↑ neutrophils and macrophages in BALF ↑ inflammatory cells infiltration and goblet cells based on H&E and PAS staining ↑ S100A9, caspase-1, IL-1 β , IL-17, IFN- γ , TNF- α and myeloperoxidase proteins in western blot analysis	45
OVA	CFA	C57BL/6	i.p.	↑ plasmacytoid dendritic cells, exudate macrophages, and B cells ↑ neutrophils in BALF and lung ↑ Th1 cytokine IFN- γ	43
	Alum	C57BL/6	i.p.	↑ interstitial macrophages and myeloid dendritic cells ↑ eosinophils in BALF and lungs ↑ IL-5 and IL-13 ↑ basophils and mast cells in lung tissue	
OVA	Alum	BALB/c	i.p.	↑ eosinophils number ↑ IL-4, IL-5, IL-13 and IL-33 in BALF Moderate inflammation (only bronchi and vessels of the lungs infiltrated with inflammatory cells)	58
	LPS	BALB/c	i.p.	↑ neutrophils number ↑ Th1 (IFN- γ) and Th17 (IL-17A) in BALF Severe inflammation (nearly whole lung infiltrated with inflammatory cells)	
HDM	Alum	BALB/c	s.c.	↑ IgE level ↑ Th2 cytokines	56
HDM	CFA	C57BL/6	s.c.	↑ macrophage MIF in BALF ↑ mixed eosinophilic/neutrophilic response AHR	47
OVA	CFA	BALB/c	s.c.	↑ neutrophils count ↑ inflammatory cell infiltration AHR	46
OVA	LPS	BALB/c	i.n.	↑ Th2 (IL-4, IL-5, IL-13) and Th17 (IL-17) ↓ Th1 (IFN- γ) and Treg (TGF- β , IL-10) ↑ GATA3, T-bet, and ROR- γ t expression ↓ T-bet, Foxp3 and IL-10 expression AHR	57

Abbreviations: i.p.: intraperitoneal; s.c.: subcutaneous; i.n.: intranasal; ILCs: innate lymphoid cells; LN: lymph node; IFN- γ : interferon-gamma; Th: T helper cells; IL: interleukin; BALF: Bronchoalveolar lavage fluid; T-bet: T-box transcription factor TBX21; ROR γ t: retinoic acid receptor-related orphan receptor gamma t; H&E: hematoxylin and eosin; PAS: periodic acid-schiff; S100A9: S100 calcium-binding protein A9; TNF- α : tumor necrosis factor alpha; IgE: immunoglobulin E; MIF: migration inhibitory factor; AHR: airway hyperresponsiveness; Treg: regulatory T cells; TGF- β : transforming growth factor-beta; Foxp3: forkhead box protein 3.

Table 2: Acute allergic airway inflammation in acute asthmatic mouse models

Strain/ge	Allergen	Sensitization/ro	Challenge/ro	Responses to challenge	References
BALB/c Female	OVA	Day 0 and 7 OVA + alum i.p.	Day 14-18 OVA i.n.	AHR and airway inflammation	59
BALB/c Female	HDM	Day 0 and 7 HDM + alum i.p.	Day 14-25 HDM i.n.	AHR, inflammatory cells infiltration, eosinophilia, Th2 cytokines and IL-33 secretion	60
BALB/c Male	OVA	Day 0 and 14 OVA + alum i.p.	Day 21-23 OVA Aerosol	Neutrophils and eosinophil infiltration airway wall thickening	52
C57BL/6 Female	OVA	Day 0 and 5 OVA + alum i.p.	Day 12 and 13 OVA Aerosol	Neutrophilia and airway inflammation	35
BALB/c Female	OVA	Day 0 and 14 OVA + alum i.p.	Day 28-30 OVA Aerosol	Inflammatory cells infiltration, Th2 cytokines secretion, eosinophilia	61
BABL/c Male	OVA	Day 7 and 14 OVA + alum i.p.	Day 21-23 OVA Aerosol	Leukocytes infiltration, eosinophilia and TNF- α , IL-1 β , IL-6, TGF- β , and IFN- γ secretion	62
Balb/c Male	OVA	Day 0, 2, 4, 7, 9 and 10 OVA i.p.	Day 15, 18 and 21 OVA i.t.	Inflammatory cells infiltration, muscle and epithelial thickening, epithelial desquamation, goblet cell metaplasia, and collagen deposition	17
BALB/c Female	OVA	Day 1 and 14: OVA + alum i.p.	Day 25-28 OVA i.n.	Inflammatory cells inflammation and IL-5 and IL-13 secretion	63
C57BL/6 Female	OVA	Day 1 and 15 OVA + alum i.p.	Day 21-23 OVA Aerosol	Aberrant miRNAs profile in the CD4 ⁺ T lymphocytes	64
BALB/c Female	OVA	Day 1, 8 and 15 OVA + alum i.p.	Day 16-22 OVA Aerosol	Airway inflammation and remodeling, inflammatory cells infiltration and Th2 cytokines secretion	65

Abbreviations: i.p.: intraperitoneal; i.n.: intranasal; i.t.: intratracheal; OVA: ovalbumin; HDM: house dust mite; alum: aluminium hydroxide; AHR: airway hyperresponsiveness; Th: T helper cells; IL: interleukin; TNF- α : tumor necrosis factor alpha; TGF- β : transforming growth factor-beta; IFN- γ : interferon-gamma; miRNAs: micro ribonucleic acids

Chronic allergen challenge model

A chronic mouse model with prolonged allergen challenges overcomes several issues encountered in the acute mouse model. Significant differences in AHR, airway remodeling, and inflammatory profiles between acute and chronic asthmatic models have been observed in clinical asthma. The chronicity of allergen exposure is a critical concern in the acute model, as the sensitization and challenge procedures may not induce persistent changes in airway inflammation, unlike in humans. Various chronic sensi-

zation and challenge protocols have been employed, with some summarized in **Table 3**.

Chronic allergen challenge contributes to persistent airway remodeling, depicted by collagen deposition, airway inflammation, goblet cell hyperplasia, and eosinophilia in the mouse model^{73,74}. The chronic model typically spans 4 to 12 weeks, starting with allergen sensitization followed by repeated low-level allergen exposure. Different types of allergens have been used to simulate the chronic model, and adjuvant-free protocols have been employed to imitate the natural sensitization that occurs in hu-

mans⁷⁵.

The presence of T cells is essential for an immediate response to recurrent allergen exposure⁷⁶. Chronic allergen exposure has demonstrated a CD4⁺ and CD8⁺ T cell-dependent effect on airway inflammatory cell infiltration and AHR^{75,77}. Moreover, eosinophils play an important role in the remodeling process by altering the structure of airway nerves, inducing AHR and fibrosis, and thereby increasing allergen sensitivity in eosinophilic asthma associated with chronic allergen exposure⁷⁸. Extensive research using chronic murine asthma models has explored the roles of some proteins, such as the WNT5A ligand⁷⁹, microRNA-221⁸⁰, and IL-33⁷³, to understand their effects on asthma pathogenesis.

The chronic mouse model has successfully replicated key features of human asthma and is currently employed to study potential therapeutic treatments applicable at the clinical stage. Extracellular vesicles derived from human umbilical cord mesenchymal stem cells have shown therapeutic potential in the chronic asthma model, particularly in a hypoxic environment⁸¹. This study demonstrated significant attenuation of airway inflammation, represented by the depletion of inflammatory cells, eosinophils, and Th2 cytokines, and amelioration of airway remodeling, accompanied by decreases in alpha-smooth muscle actin (α -SMA), collagen type 1, and transforming growth factor-beta (TGF- β) 1 signaling pathway expression.

Additionally, this model is also used to gain a better understanding of biochemical changes within complex tissue samples of potential anti-asthmatic compounds⁸². Novel imaging techniques that combine the analytical approaches of focal plane array (FPA) and synchrotron Fourier-transform infrared (S-FTIR) enable the investigation of broader molecular changes surrounding the airways and identification of types of collagen deposition present in the chronic asthma model, further supporting the analysis of conventional methods.

However, several hindrances related to the chronic mouse model were identified when compared to human asthma. In humans, asthma often develops spontaneously in early life alongside immature lung development, compared to the fully developed lungs of mice at birth, necessitating artificial allergen and adjuvant sensitization⁸³. The route, amount, and frequency of allergen exposure in controlled conditions of allergic airway mouse models differ from the natural and acquired immune responses of asthma exacerbation in humans and do not reflect patient heterogeneity^{84,85}.

Moreover, the extended period of inhaled antigen exposure in mice induces tolerance, described by changes in inflammatory cell profiles, airway inflammation, and AHR, limiting the opportunities to investigate the chronic model and the underlying pathways^{86,87}. Nevertheless, allergen tolerance provides some advantages for studying the effect of certain parameters associated with asthma for therapeutic development. Inhaled allergens may induce an inappropriate Th2-cell inflammatory response, and this adverse reaction can be obscured via the local inhalation tolerance process to restore airway homeostasis⁸⁸ and regulation of free IgE⁸⁹, thereby diminishing asthma symptoms.

While invaluable for research, these chronic mouse models pose significant ethical challenges. Prolonged suffering, due to repeated allergen exposure leading to chronic inflammation, AHR, and airway remodeling, can cause discomfort, breathing difficulties, and potentially pain over extended periods⁹⁰. Assessing pain and distress can be challenging, as subtle behavioral changes may indicate underlying suffering but are difficult to interpret definitively⁹¹. Therefore, researchers must carefully optimize research protocols by minimizing the duration and intensity of allergen exposure, balancing the need to reduce distress with the requirement to obtain meaningful data. Animals should also be monitored regularly for signs of distress, including routine assessment of respiratory function and behavior.

FUTURE PERSPECTIVES AND CONCLUSION

Allergen sensitization and challenge in mouse models represent classical protocols for manifesting asthma pathophysiology. Researchers are striving to model specific disease phenotypes that accurately replicate the complex nature of human asthma. While acute allergen challenges effectively represent several hallmarks of asthma, they fall short of capturing certain features of chronic asthma. Therefore, the development of chronic allergen challenge models aims to deepen understanding of disease mechanisms and discover novel therapeutic potentials.

Allergic mouse models require active sensitization, typically introduced with adjuvants administered intraperitoneally or subcutaneously alongside the allergen. These methods are less intrusive and do not require sedation, but they may induce tolerance. As a result, models without adjuvants have been developed to induce sensitization in the airways via intranasal instillation, simulating the natural exposure of humans to airborne allergens. This model has

Table 3: Chronic allergic airway inflammation in chronic asthmatic mouse models

Strain/gen	Allergen	Sensitization/ro	Challenge/route	Response to challenge	References
BALB/c Female	OVA	Day 0, 7, and 14 OVA + alum i.p.	Day 21-55 OVA Aerosol/i.n.	Airway remodeling, inflammatory cells infiltration, eosinophilia, increased mucus production and IL-4 and IL-13 secretion	82
BALB/c Female	OVA	Day 0, 7, 14, and 21 OVA + alum s.c.	Days 33 and 35: OVA i.n.	AHR, airway inflammation, and remodeling, Th2 cytokines, TSLP, IL-33 and IL-25 secretion, goblet cell hyperplasia, increased TNF- α , and collagen deposition	92
BALB/c Female	OVA	Day 0, 14, 28 and 42 OVA + alum i.p.	Day 21-42 OVA Aerosol	Airway remodeling, inflammatory cells infiltration, elevated IgE, IL-6 and IL-13	93
Balb/c Female	HDM	Days 0, 7, and 14 HDM i.p.	Day 21-28 HDM i.n.	Inflammatory cells infiltration, Th2 cytokines secretion and specific IgE production, airway wall thickening, mucosal metaplasia, collagen deposition, goblet cell hyperplasia and mucus hypersecretion.	30
BALB/c Female	OVA	Day 1 and 14 OVA + alum i.p.	Day 14, 17, 21, 24, 27, 60, 69, 71, 73, 74, and 75 OVA i.n.	Inflammatory cells infiltration, Th2 cytokine, IL-17, TNF- α and high mobility group box protein 1 secretion.	94
BALB/c Female	OVA	Day 1, 2 and 3 OVA + alum i.p.	Day 14, 17, 21, 24, 27, 60, 69, 71, 73, 74, and 75 OVA i.n.	Airway remodeling, inflammatory cells infiltration and Th2, Th1, IL-17 and IL-22 cytokines secretion and collagen deposition	95
BALB/c Female	OVA	Day 1 and 14 OVA + alum i.p.	Day 28, 30, 32, 34, 36, 38, 40, 42 and 44 OVA Aerosol	Airway inflammation, fibrotic airway remodeling and inflammatory cells infiltration	63
C57BL/6 Female	HDM	Day 1 HDM i.n.	Day 2-36: HDM i.n.	Th2-mediated eosinophilic inflammation and IL-12 and IL-6 production	96
Balb/c Male	OVA	Day 0 and 14 OVA+ alum i.p.	Three times per week for 9 weeks OVA Aerosol	Day 87: AHR, inflammatory cells infiltration, eosinophilia, and mucus hypersecretion	97
C57BL/6 Female	HDM	Day 0 and 7: HDM i.n.	five times per week for three weeks, rested (4-8 week) and rechallenged HDM i.n.	24 hours after the final challenge: AHR, increased CD4 ⁺ T cells and dendritic cells	98

Abbreviations: i.p.: intraperitoneal; i.n.: intranasal; s.c.: subcutaneous; OVA: ovalbumin; alum: aluminum hydroxide; IL: interleukin; Th: T helper cells; TSLP: thymic stromal lymphopoietin; TNF- α : tumor necrosis factor alpha; IgE: immunoglobulin E; HDM: house dust mite; AHR: airway hyperresponsiveness.

proven effective in producing a phenotype of asthma comparable to that of the traditional adjuvant model. In allergen challenge, aerosol and intranasal routes are likely closer to mimicking human exposure than the intratracheal approach. The allergen OVA may inadvertently induce tolerance with repeated and prolonged exposure, in contrast to HDM, which exhibits persistent airway inflammation, making it more suitable for modeling chronic asthma. Therefore, adjuvant-free models and aeroallergen exposure may be more relevant in mimicking human asthma for the development of new treatments and preventive approaches. Despite the shortcomings of both acute and chronic allergic asthma models, ongoing research aims to improve protocols to enhance our understanding of asthma at the cellular and molecular levels.

ABBREVIATIONS

α -SMA (Alpha-smooth muscle actin), AHR (Airway hyperresponsiveness), Alum (Aluminum hydroxide), APC(s) (Antigen-presenting cell(s)), BALF (Bronchoalveolar lavage fluid), CFA (Complete Freund's Adjuvant), ec (Epicutaneous), Foxp3 (Forkhead box protein 3), FPA (Focal plane array, an imaging technique), H&E (Hematoxylin and eosin staining), HDM (House dust mite), IFN- γ (Interferon-gamma), IgE (Immunoglobulin E), IL (IL-4, IL-5, IL-13, IL-17, etc.) (Interleukin), ILC(s) (ILC1, ILC2, ILC3) (Innate lymphoid cell(s)), i.n. (Intranasal), i.p. (Intraperitoneal), i.t. (Intratracheal), LN (Lymph node), LPS (Lipopolysaccharide), MHC (Major histocompatibility complex), miRNAs (Micro ribonucleic acids), OVA (Ovalbumin), PAS (Periodic acid-Schiff staining), ROR γ t (Retinoic acid receptor-related orphan receptor gamma t), s.c. (Subcutaneous), S-FTIR (Synchrotron Fourier-transform infrared spectroscopy), S100A9 (S100 calcium-binding protein A9), T-bet (T-box transcription factor TBX21), TGF- β (Transforming growth factor-beta), Th (Th1, Th2, Th17) (T helper cells), TLR4 (Toll-like receptor 4), TNF- α (Tumor necrosis factor alpha), and TSLP (Thymic stromal lymphopoietin)

ACKNOWLEDGMENTS

None.

AUTHOR'S CONTRIBUTIONS

Bushra Solehah Mohd Rosdan served as the primary author and was responsible for the initial drafting and subsequent editing of the manuscript. Nurul Asma Abdullah provided supervision, conducted

critical reviews, and contributed to manuscript revisions. All authors have read and approved the final version of the manuscript.

FUNDING

This study was funded by Research University Grant (1001/PPSK/8012344) from Universiti Sains Malaysia.

AVAILABILITY OF DATA AND MATERIALS

Not applicable.

ETHICS APPROVAL AND CONSENT TO PARTICIPATE

Not applicable.

CONSENT FOR PUBLICATION

Not applicable.

COMPETING INTERESTS

The authors declare that they have no competing interests.

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