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The Complicated Role of Nuclear Factor Erythroid-Derived 2-Like 2 in Allergy and Asthma

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ABSTRACT

Nuclear factor erythroid-derived 2-like 2 (Nrf2) is a stress-activated transcription factor that is highly responsive to oxidative stress and electrophilic stimuli. Upon activation, Nrf2 upregulates a battery of cytoprotective genes meant to prevent cell death or damage. In many models of inflammation, Nrf2 protects against the immune response and decreases injury, including in the context of asthma and allergy. However, in some models of asthma and allergy, Nrf2 either does not play a role or can even exacerbate inflammation. In general, the reasons behind these discrepancies are not clear and the mechanisms by which Nrf2 modulates immune response are largely uncharacterized. The aim of this review is to highlight current literature assessing the role of Nrf2 in

allergy and asthma to understand Nrf2 as a potential therapeutic target.

SIGNIFICANCE STATEMENT

Nuclear factor erythroid-derived 2-like 2 (Nrf2) is an important immune mediator that modulates numerous immune cell types in various inflammatory diseases, including allergy and asthma. There is considerable interest in Nrf2 as a drug target in inflammation, which is complicated by the complex nature of Nrf2 in the immune system. This review focuses on the role of Nrf2 in asthma and allergy, including in regulating immune cell function and in detoxifying xenobiotics that exacerbate these diseases.

Introduction

Nuclear factor, erythroid-derived 2, -like 2 (Nrf2; encoded by Nfe2l2) was identified as a member of the Cap “n” Collar basic leucine zipper transcription factor family in 1994 (Moi et al., 1994). It was first isolated from K562 cells (a human chronic myelogenous leukemia cell line) where it was found binding to the β -globin locus control region. The name Nrf2 was adopted from the nomenclature for the transcription factor known as nuclear factor erythroid-derived 2-like 1 (Nrf1), which was named because of its similarities with nuclear factor erythroid 2 (NF-E2) (Chan et al., 1993). However, unlike NF-E2, which is essential for erythropoiesis and is neonatally lethal in mice when knocked out, Nrf2 gene function is not necessary for blood cell differentiation, and homozygous

Nrf2-null offspring are fertile and produce normal litter sizes (Shivdasani and Orkin, 1995; Chan et al., 1996; Williams et al., 2016). Subsequent research has demonstrated that Nrf2 forms a heterodimer with small musculoaponeurotic fibrosarcoma (Maf) proteins and regulates the expression of Nrf2-target genes involved in oxidative stress response and drug detoxification such as NAD(P)H quinone-oxidoreductase-1 (NQO1), γ -glutamyl cysteinyl synthetase, and heme-oxygenase-1 (HO-1) by binding to the antioxidant response elements (AREs) in the promoter regions of these genes (Itoh et al., 1997; Thimmulappa et al., 2002; Nguyen et al., 2003; Tong et al., 2006). In addition to its antioxidant and detoxification roles, Nrf2 plays a protective anti-inflammatory role in many different animal models of inflammation. Although much is known about the mechanism by which Nrf2 protects against reactive toxicants and oxidative stress, the mechanism by which Nrf2 modulates immune cell function is not nearly as well understood. We recently published a review focused on the role of Nrf2 in autoimmunity and infectious disease (Freeborn and Rockwell, 2021). In addition, others have reviewed the role of Nrf2 in inflammation and in crosstalk with nuclear factor kappa B (NF- κ B) (Li et al., 2008a; Wardyn et al., 2015; Ahmed

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ABBREVIATIONS: ACD, allergic contact dermatitis; AD, atopic dermatitis; AhR, aryl hydrocarbon receptor; ARE, antioxidant response element; BALF, bronchoalveolar lavage fluid; BMDC, bone marrow-derived dendritic cells; CYP2E1, cytochrome P450 family 2 subfamily E member 1; DC, dendritic cells; DEP, diesel exhaust particles; DMF, dimethyl fumarate; DNCB, 2,4-dinitrochlorobenzene; ECH, enoyl-CoA hydratase; H O-1, heme oxygenase 1; Keap1, Kelch ECH-associating protein 1; KO, knockout; Maf, musculoaponeurotic fibrosarcoma; miR, microRNA; Neh, Nrf2-ECH homology; NF-E2, nuclear factor,erythroid 2; NQO1, NAD(P)H quinone-oxi-doreductase-1; Nrf2, nuclear factor,erythroid-derived 2,-like 2; OVA, ovalbumin; PM, particulate matter; PMN, polymorphonuclear neutrophil; UFP, ultrafine particle; WT, wild-type.

et al., 2017; Mohan and Gupta, 2018; Staurengo-Ferrari et al., 2019; Saha et al., 2020; Bhandari et al., 2021; Freeborn and Rockwell, 2021). To complement these previous reviews, the purpose of the present paper is to review what is known about the role of Nrf2 in asthma and allergy.

Brief Historical Perspective

Structure and Regulation of Nrf2 Protein. The Nrf2 protein comprises six highly conserved Nrf2-erythrin-CoA hydratase (ECH) homology (Neh) domains (Neh1–6) (Ma, 2013). Each domain has a distinct function. Neh1 contains the Cap “n” Collar-basic leucine zipper domain, which allows Nrf2 to heterodimerize with small Mafs (MafF/G/K) and to bind to the ARE (Satta et al., 2017). The Neh3, Neh4, and Neh5 domains contribute to Nrf2 transactivation (Satta et al., 2017). The redox-insensitive Neh6 domain is essential for the degradation of the Nrf2 in stressed cells (McMahon et al., 2004). In contrast, in homeostatic cells, the Glu₇₉-Thr₈₀-Gly₈₁-Glu₈₂ (ETGE) tetrapeptide motif and Asp₂₉-Leu₃₀-Gly₃₁ (DLG) motif within the redox-sensitive Neh2 domain interact with Kelch ECH-associating protein 1 (Keap1). Keap1 represses Nrf2 by associating with a ubiquitin E3 ligase that polyubiquitinates the Nrf2 protein, resulting in its subsequent degradation by the 26S proteasome (Canning et al., 2015). However, under conditions of oxidative stress, or in the presence of electrophilic xenobiotics, the conformation of Keap1 is changed, which disrupts the polyubiquitination and degradation of Nrf2. Subsequently, Nrf2 translocates and accumulates in the nucleus, where it activates Nrf2-target genes to induce expression of antioxidant proteins, detoxification enzymes, and other protective genes (Itoh et al., 2003; Kensler et al., 2007; Ma, 2013; Ahmed et al., 2017; Yamamoto et al., 2018).

The regulation of the Nrf2 signaling pathway is complex. Nrf2 stability can be enhanced through inhibition of Keap1 or disruption of the Nrf2-Keap1 interaction (Kansanen et al., 2013). Nrf2 activators, such as electrophilic compounds, can hinder the degradation of Nrf2 by modifying key cysteines of Keap1 and thereby changing the conformation of Keap1. Proteins that interact or compete with Keap1 can disrupt the Nrf2-Keap1 association, resulting in the accumulation of newly synthesized Nrf2. Nrf2 can also be regulated at transcription level. The promoter region of Nfe2l2 (the gene that encodes Nrf2) contains xenobiotic response element-like (XRE) sequences, which are the binding sites of the aryl hydrocarbon receptor (AhR) (Miao et al., 2005). AhR is a basic helix-loop-helix transcription factor regulating xenobiotic metabolism and works in close concert with the transcription factor Nrf2 (Miao et al., 2005; Yeager et al., 2009). Published studies indicate that the Nfe2l2 gene is directly upregulated by AhR activation, and conversely, Nrf2 induces AhR expression and thus triggers several downstream events of the AhR signaling cascade (Li et al., 2019). Thus, these two xenobiotic sensors induce one another. In contrast to AhR, the NF- κ B p65 subunit represses the Nrf2-ARE pathway at the transcriptional level (Liu et al., 2008). Thus, Nrf2 expression and activity are modulated by multiple other mediators in a complex network.

Beyond the classic Nrf2/Keap1-mediated activation and degradation and transcriptional regulation of Nrf2, Nrf2 can also be regulated at the post-transcriptional level. Numerous microRNA (miR) molecules can repress Nrf2 expression by sequence-specific binding, particularly in the context of cancer (Tonelli et al., 2018). miR-29-b1 and miR-144 downregulate Nrf2 expression directly within the cytoplasm (Ayerdemonstrated elevated type 2 cytokines such et al., 2015). In contrast, microRNA targeting Keap1, such as miR-200a and miR7, represses Keap1 expression corresponding with Nrf2 nuclear translocation and activation (Eades et al., 2011; Kabaria et al., 2015).

Evidence that Nrf2 Modulates the Immune System: The Development of Autoimmunity in Nrf2-Knockout Mice. Considerable evidence points to an important role for Nrf2 in regulating the immune system. Some of the strongest and earliest indications that Nrf2 regulates immunity were in the area of autoimmunity. Between 2004 and 2006, two different groups described the spontaneous development of an autoimmune disease in Nrf2-knockout (KO) mice that resembled systemic lupus erythematosus. The disease was female-predominant and characterized by autoantibody production (anti-dsDNA), formation of antibody complexes in kidney, and development of glomerulonephritis that resulted in kidney injury, among other pathologies in these animals (Li et al., 2004a; Ma et al., 2006). In addition to the spontaneous development of autoimmunity, many animal models of autoimmune disease were found to be more severe in Nrf2-KO mice. Antibody-induced rheumatoid arthritis (RA) is exacerbated in Nrf2-null mice, with an increase in joint damage and a decrease in the expression of antioxidant genes, such as HO-1, γ -glutamyl cysteinyl synthetase, and thioredoxin (Wruck et al., 2011). In experimental autoimmune encephalomyelitis (EAE), Nrf2-null mice develop symptoms earlier and have increased severity of clinical outcomes when compared with wild-type (WT) controls (Johnson et al., 2010; Larabee et al., 2016). Consistent with these animal studies, the Nrf2 activator dimethyl fumarate (DMF) is used clinically to treat multiple sclerosis (Tecfidera) and psoriasis (Skilarence) in humans (Bomprezzi, 2015). Furthermore, recent studies using patient samples and animal models indicate that DMF may be beneficial for the treatment of systemic sclerosis patients as well (Toyama et al., 2018; Kourakis et al., 2020). There is evidence to show that several other Nrf2 activators, such as 3H-1,2-dithiole-3-thione, sulforaphane, dimethyl itaconate, and A-1396076, can also ameliorate autoimmune-mediated inflammation in rodents under experimental conditions (Geisel et al., 2014; Kuo et al., 2016, 2020; Goess et al., 2020). The therapeutic potential of Nrf2 activators in the treatment of autoimmune disease remains an active area of research that has yielded many developments over the years. We have recently written a comprehensive review of research in this area (Freeborn and Rockwell, 2021). Taken together, these studies provide strong evidence that the Nrf2 signaling pathway modulates immune cell function and paved the way for an explosion of research into the role of Nrf2 in the immune system, including in atopic diseases. The present manuscript reviews the role of Nrf2 in asthma and allergy. In an effort to keep the review focused, we have largely restricted the studies with animal models to those where a causative role for Nrf2 has been established (with a few exceptions for disease models with fewer studies published).

Key Recent Advantages

The Role of Nrf2 in Asthma and Airway Allergic Inflammation. Atopic diseases, such as asthma, allergic rhinitis, skin allergy, and food allergy, are the sixth leading cause of chronic illness in the United States (<https://acaai.org/news/allergy-facts/>). Asthma is a condition in which airways are hyperreactive to irritants and allergens, leading to uncontrolled airway constriction that can result in hypoxia and death (Barnes, 2018). Allergic lung inflammation and asthma are lower respiratory tract disorders characterized by reversible airflow obstruction, as well as wheezing, coughing, and chest tightness upon exposure to allergens, such as dust mites, cockroaches, mold, pollen, and animal dander (Bousquet et al., 2000). Allergic rhinitis is an upper airway disorder characterized by nasal itching, sneezing, and nasal obstruction caused by inflammation of the nose upon contact of the nasal mucosa with certain allergens, such as pollen, mold, diesel fumes, and air pollutants (Pawankar et al., 2011; Varshney and Varshney, 2015; <https://acaai.org/allergies/allergic-conditions/hay-fever/>).

TABLE 1
Studies focused on the role of Nrf2 in allergic airway and asthma

Disease Model	Environmental Factors/Activators	Genetic Model	Effect of Knockout/ Knockdown (Nrf2 or Keap1)	References
OVA-induced allergic airway inflammation/asthma		CD1: ICRWT & Nrf2(−/−)	↑ AHR, ↑ inflammatory cell infiltrate in lung, ↑ lipid peroxidation, ↑ IL-4, IL-13, ↑ mucus cell metaplasia	(Rangasamy et al., 2005)
	Diesel	C57BL/6WT & Nrf2(−/−)	↑ AHR, ↑ inflammatory cells infiltrate in lung, ↑ IL-5, ↑ mucus cell hyperplasia	(Li et al., 2010)
	ambient UFP (<0.18 μm)	BALB/c WT & Nrf2(−/−), DC from WT & Nrf2(−/−)	↑ Adjuvant effect of intranasally instilled UFP, ↑ eosinophil count and IL-13 in BAL, V serum IgG1	(Li et al., 2013)
		<i>Tam-Keap1</i> ^{−/−}	↓ AHR, ↓ eosinophilic inflammation, and ↓ IL-4 and IL-13 in BAL	(Sussan et al., 2015)
Ragweed Extract-induced asthma		BMDC and lung DC from WT Nrf2 (−/−)	↑ CD80, CD86, and MHCII on DC, ↑ IL-6 and TNF-α ↑	(Rangasamy et al., 2010)
Cl ₂ -induced inflammation and airway hyperresponsiveness		BALB/cWT & Nrf2(−/−)	↓ mRNA for antioxidant genes (NQO1 and GPX2)	(Ano et al., 2017)
IL-33-induced allergic lung inflammation PM2.5-induced airway inflammation	CDDO-Im, 15d-PGJ ₂	C57BL/6WT & Nrf2(−/−)	Transient ↑ in IL-33, ↑ ILC2 proliferation	(Nagashima et al., 2019)
		C57BL/6WT & Nrf2(−/−)	↓ Inflammatory infiltrate, ↓ oxidative stress, ↓ lung injury, ↓ CYP2E1	(Ding et al., 2021)
		C57BL/6WT & Nrf2(−/−)	↑ Lung function, ↓ alveolar wall thickening, ↓ in low-grade inflammation	(Jiang et al., 2021)

CDDO-Im, 1-[2-cyano-3-,12-dioxooleana-1,9(11)-dien-28-oyl] imidazole; GPX2, glutathione peroxidase 2; ILC2, type 2 innate lymphoid cells; 15d-PGJ₂, 15-deoxy-prostaglandin J₂.

Oxidative stress in airway cells has been proposed to be a contributing factor in the pathogenesis of hyperreactive airway disorders. Extensive studies, listed in Table 1, have been conducted on the role of Nrf2 in allergic airway inflammation and asthma. For instance, genetic deletion of Nrf2 in mice increased airway inflammation and hyperresponsiveness to ovalbumin (OVA) (Rangasamy et al., 2005). Specifically, the authors demonstrated elevated type 2 cytokines such as interleukin (IL)-4 and IL-13 in bronchoalveolar lavage fluid and splenocytes in Nrf2-deficient mice after OVA challenge (Rangasamy et al., 2005). This group further showed that Nrf2 is specifically cytoprotective in airway epithelial cells in this model (Sussan et al., 2015). Consistent with this, Hirai et al. (2019) showed that low Nrf2 expression correlated with increased severity in human asthma patients. A study by Brown et al. (2015) showed that consumption of the Nrf2 activator sulforaphane ameliorated bronchoconstriction in 60% of individuals with moderate asthma. Interestingly, sulforaphane administration was associated with a decrease in Nrf2 target gene expression in these patients. However, 20% of the patients who consumed sulforaphane experienced exacerbated bronchoconstriction, suggesting that sulforaphane was not protective in all subsets (Brown et al., 2015).

Similarly, a protective role of Nrf2 has been implied in chemical-induced asthma. Chlorine gas-induced airway inflammation and asthma were significantly greater at 48 hours postexposure in Nrf2-deficient mice compared with WT (Ano et al., 2017). Other studies have directly shown the therapeutic effect of Nrf2 activation in acute allergic asthma models. Both genetic activation of Nrf2 via deletion of its suppressor protein Keap1 and pharmacological activation of Nrf2 via 2-trifluoromethyl-2'-methoxychalcone improved the cytoprotective function of the airway epithelium in an OVA-induced asthmatic mouse model (Sussan et al., 2015).

To prevent asthma-induced bronchoconstriction, long-acting beta agonists and inhaled steroids are chiefly used. Steroids, such as dexamethasone, act as a local anti-inflammatory agent and preserve the integrity of the airway epithelium, preventing the airway irritation that leads to constriction. However, steroids can lose their effectiveness over time. Activation of Nrf2 restores steroid sensitivity in a mouse model of asthma (Sakurai et al., 2018). The protective effects of Nrf2 in these models are likely mediated by Nrf2 target genes rather than Nrf2 itself. Aldehyde oxidase is a well described target gene of Nrf2 and is instrumental in the formation of tight junctions and adherent junctions in airway epithelium (Shintani et al., 2015). Genetic ablation of Nrf2 in cell lines led to decreased signaling via the aldehyde oxidase pathway (Shintani et al., 2015). Overall, the evidence points to a protective role for Nrf2 in this model of asthma, which is likely mediated at least in part by maintaining the integrity of the airway epithelial barrier.

House dust mites (HDMs) are some of the most common perennial sources of allergens that induce asthma and allergic airway inflammation. Investigators have shown that exposure to house dust mites decreases Nrf2 levels, suggesting that antioxidant and anti-inflammatory pathways are inhibited, which may contribute to the dysregulation of sinonasal epithelial cell barrier function and the development of asthma (London et al., 2017). Nrf2 deficiency has also been shown to exacerbate the response of lung dendritic cells to ragweed extract. Specifically, Nrf2-deficient lung dendritic cells cultured in vitro showed greater induction of TNF-α and IL-6 in response to ragweed extract (Rangasamy et al., 2010). OVA is another major allergen that induces allergic reactions, particularly in experimental animal models of allergy and asthma. Several studies have shown that OVA induces several hallmarks of allergic reactions, including sinonasal inflammation and most prominently, asthma (Rangasamy et al., 2005; Sussan et al., 2015).

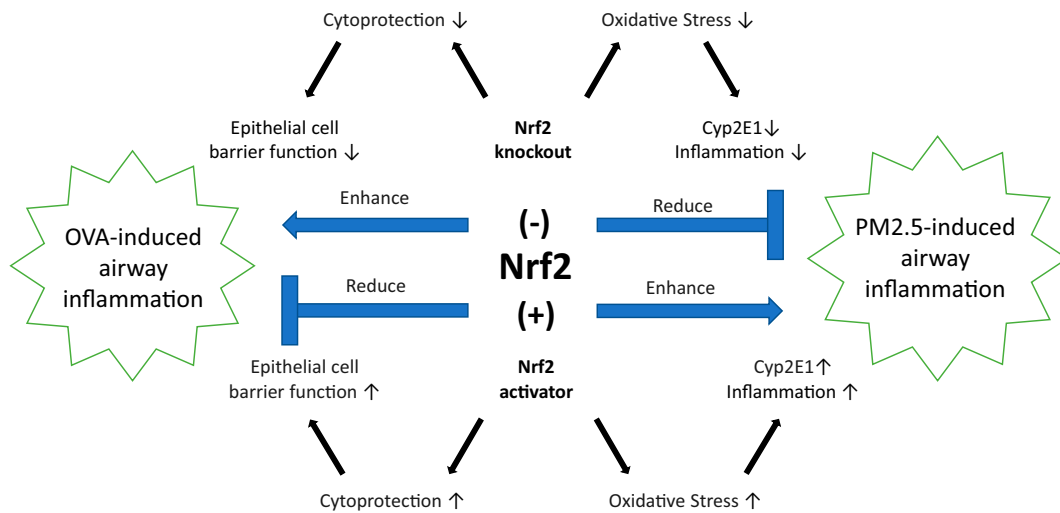


Fig. 1. The role of Nrf2 in airway inflammation. Nrf2 can either contribute to or be protective against airway inflammation. Nrf2 has been shown to be protective in OVA-induced airway inflammation by promoting airway epithelial cell barrier function. In contrast, Nrf2 contributes to airway inflammation caused by PM_{2.5} through a mechanism that is thought to involve CYP2E1.

Nrf2 deficiency is associated with mucus cell hyperplasia and eosinophilic lung infiltration in a murine OVA-induced asthma model. In these studies, bronchoalveolar lavage fluid (BALF) and splenocytes of Nrf2-deficient mice showed increased expression of IL-4 and IL-13 after OVA exposure (Rangasamy et al., 2005). Other studies have examined the role of Nrf2 in the airway epithelium after sensitization to OVA, in which they demonstrated that genetic and pharmacological Nrf2 activation suppressed asthma (Sussan et al., 2015). The suppression of OVA-induced asthma by Nrf2 was associated with increased expression of the Nrf2 target gene NQO1 and a significant decrease in infiltrating neutrophils and lymphocytes. The secretion of cytokines like IL-4 and IL-13 in BALF was also decreased (Sussan et al., 2015). In contrast to these studies, a study by Seumois et al. (2014) in asthmatic and non-asthmatic individuals suggests that Nrf2 is a strong positive determinant of Th2 differentiation in CD4 T cells in humans. The data show significant enrichment of Nrf2 binding sites in Th2-specific enhancers. Consistent with this, our group has shown that activation of Nrf2 promotes Th2 differentiation of isolated murine CD4 T cells (Rockwell et al., 2012).

Epidemiologic data have shown that there is a link between certain genetic polymorphisms in the *N*-acetyltransferase 2 (NAT2), glutathione S-transferase Pi 1 (GSTP1), and *Nfe2l2* genes and the occurrence of asthma in children if they are exposed to acetaminophen. They also reported an increase in serum eosinophils when children with a specific Nrf2 polymorphism were taking acetaminophen (Kang et al., 2013). However, it is not clear whether the impact of the Nrf2 polymorphism in this study is due to effects on immune cell function, detoxification of acetaminophen, or some combination of these.

There has been interest in Nrf2 as a therapeutic target for chronic respiratory illnesses, including asthma (Wang et al., 2018). For example, targeted activation of Nrf2 in female mice suppresses allergic lung inflammation and alleviates OVA-induced asthma (Nagashima et al., 2019). Nrf2 activation decreases type 2 innate lymphoid cells (ILC2s), which release the major allergic airway inflammation-inducing cytokines IL-4 and IL-13 (Nagashima et al., 2019). Recently, it has been shown that Nrf2 activation by alpha lipoic acid also suppressed cytokines TNF- α and IL-6, which are associated with non-Th2-mediated airway inflammation (Van Nguyen et al., 2020).

Nrf2, Air Pollution, and Asthma/Allergy. Air pollutants and diesel fumes are among the common causes of allergic rhinitis (Li et al.,

2020; Jung et al., 2021). Air pollutants, such as carbon monoxide, nitrogen dioxide, lead, ozone, and other particulate matter (PM), contribute to oxidative stress and inflammation in rhinitis (Lodovici and Bigagli, 2011; Pardo et al., 2020). Diesel fumes are one of the major contributors to air pollution, consisting of particulate matter composed of metals and polycyclic aromatic hydrocarbons. These diesel-derived particles have an aerodynamic diameter of 2.5 μ m or smaller (PM_{2.5}), allowing them to effortlessly penetrate the lungs. Studies have shown that exposure to PM_{2.5} induces oxidative stress by increasing cellular reactive oxygen species and inflammation, as marked by increased IL-8 expression (Buttrick et al., 2018). Studies have also indicated that exposure to PM_{2.5} from biodiesel results in increased inflammation marked by an increase in macrophage-derived TNF- α expression, as well as increased expression of Nrf2 and HO-1 proteins, suggesting that PM_{2.5} from biodiesel may stimulate the activation of Nrf2/HO-1 pathways (Cattani-Cavalieri et al., 2019). Exposure to PM_{2.5} from biodiesel increased expression of p-NF- κ B, which stimulates proinflammatory cytokines, such as TNF- α (Cattani-Cavalieri et al., 2019). Similarly, diesel exhaust particles (DEP) have also been shown to activate Nrf2 in macrophage and bronchial epithelial cell lines (Li et al., 2004b). Furthermore, exposure to DEP or exposure to DEP in combination with OVA sensitization resulted in increased airway hyperresponsiveness and greater immune cell counts and cytokine levels in bronchial alveolar lavage fluid from Nrf2-KO mice compared with WT mice (Li et al., 2008b, 2010). Studies with ultrafine particles (UFP) indicate that adoptive transfer of Nrf2-deficient dendritic cells treated with UFP and OVA into WT mice resulted in greater immune cell infiltrate, OVA-IgG₁, and IL-13 in the BALF after in vivo OVA sensitization compared with mice receiving similarly treated WT dendritic cells (Li et al., 2013). Taken together, these studies indicate that particulate matter can activate Nrf2 in the lung, which can have a protective effect in these models.

Although most studies indicate that Nrf2 deficiency results in worsened oxidative stress, other studies show that Nrf2 may also contribute to injury or have no effect under certain conditions (Fig. 1). A recent study using a real-ambient air exposure system demonstrated that chronic exposure to real ambient air containing high levels of PM_{2.5} caused an increase in inflammatory infiltrate, oxidative stress, and lung injury in WT, but not Nrf2-null, mice (Ding et al., 2021). The authors hypothesize that the diminished inflammation and damage observed in the Nrf2-KO mice may be due to decreased expression of CYP2E1

TABLE 2
Studies focused on the role of Nrf2 in skin allergy

Disease	Model	Effect of Nrf2-Knockout	References
Atopic Dermatitis	BALB/cWT & Nrf2 (-/-)	↓ Ear thickness; ↓ effector cell infiltration, ↓ type 2 cytokine, ↓ IgE response	(Ogawa et al., 2020b)
Contact Dermatitis	<i>Ex vivo</i> : BMDC Nrf2 (+/+) versus (-/-)	↓ Expression of antioxidant genes	(Mussotter et al., 2016)
	C57BL/6WT & Nrf2(-/-)	↑ PMN recruitment to skin during sensitization phase	(Helou et al., 2019)
	C57BL/6WT & Nrf2(-/-)	↑ Ear thickness	(El Ali et al., 2013; Helou et al., 2019)

IgE, immunoglobulin E.

(cytochrome P450 family 2 subfamily E member 1), which they postulate produces a toxic metabolite of PM_{2.5}. Further, they assert that induction of protective phase II enzymes is not sustained with long-term Nrf2 activation in their system. The diminished injury by chronic PM_{2.5} exposure in Nrf2-KO mice was corroborated in another study, where it was associated with increased p62 (sequestosome) expression and impaired intrinsic autophagy (Jiang et al., 2021). Collectively, these studies indicate that Nrf2 may contribute to lung injury resulting from chronic exposure to particulate matter under certain circumstances.

Nrf2 and Atopic Dermatitis. Atopic dermatitis (AD), also known as atopic eczema, is the most common inflammatory skin disease that is characterized as persistent eczematous lesions, itch, and discomfort. The pathophysiology of AD involves both genetics and environmental factors that can cause epidermal barrier abnormalities and inflammation of the skin from T cells (Weidinger et al., 2018).

Many studies have identified compounds with anti-inflammatory properties in both *in vitro* and *in vivo* models of AD with evidence for activation of Nrf2; however, it is important to note that a causative role for Nrf2 has not yet been established in most of these studies (Table 2). Specifically, studies have shown that sulforaphane, saponins derived from *Platycodon grandiflorum*, quercetin, macakurzcin-C derivative, and a chrysin derivative all have protective effects in animal models of atopic dermatitis, with data suggesting that these effects correlate with Nrf2 activation (Wu et al., 2019; Choi et al., 2014; Karuppagounder et al., 2015; Akram et al., 2016; Yu et al., 2019). Additional studies will be needed to identify the role of Nrf2 in these effects.

Whereas most of the evidence for a role of Nrf2 in AD comes from studies utilizing Nrf2 activators, there has been at least one study that employed Nrf2-knockout mice (Table 2). Ogawa et al. (2020b) demonstrated that expression of Nrf2 is required for the development of atopy in experimental AD induced by chronic TNCB (2,4,6-trinitro-1-chlorobenzene) treatment. Specifically, the authors showed decreased ear thickness and diminished infiltration of a number of effector cells, including mast cells, basophils, and CD4 T cells in Nrf2-null mice. These effects correlated with a marked decrease in the expression of type 2 cytokines and serum IgE levels. Overall, this study suggests a role for Nrf2 in promoting type 2 immune response, inflammation, and atopy in AD (Ogawa et al., 2020b).

The Role of Nrf2 in Contact Dermatitis. Allergic contact dermatitis (ACD) is one of the most common skin allergies in western countries, with allergens differing based on an individual's environment and personal habits (Thyssen et al., 2007). ACD is a T cell-mediated response induced by epicutaneous sensitization with an allergen that results in immunologic memory and increased inflammation at the site of re-exposure to the allergen (Vocanson et al., 2009). In contrast to atopic dermatitis, contact dermatitis is not typically associated with a pronounced type 2 immune response.

Nrf2 is thought to play a protective role in contact dermatitis through the upregulation of antioxidant and cytoprotective genes (Ma, 2013). Numerous studies have reported contact sensitizers to induce the Nrf2 pathway, and these are listed in Table 2. Cluster of differentiation (CD)34-derived dendritic cells (DC) and the THP-1 myeloid cell line were treated with various sensitizers, and an increased expression of HO-1 and NQO1 mRNA was observed. Additionally, Nrf2 protein accumulation was found only in cells treated with chemical sensitizers (Ade et al., 2009). In keratinocytes, the antioxidant and Nrf2 target gene HO-1 was significantly upregulated when exposed to contact sensitizers (Vandebriel et al., 2010). Others have found activation of Nrf2 to alleviate contact dermatitis through several compounds such as melatonin, JNJ777120 (histamine H4 receptor antagonist), 2',3'-dihydroxy-4',6'-dime-thoxychalcone, and 6-shogaol (active compound of ginger) (Park et al., 2016; Shen et al., 2017; Mohamed et al., 2018; Takada-Takatori et al., 2019). Although these studies suggest a potential protective effect of Nrf2 activators on allergic contact dermatitis, further studies with Nrf2-deficient models are needed to identify the role of Nrf2 in these effects.

To specifically address the role of Nrf2 in contact dermatitis, a study was performed using WT and Nrf2-deficient mice. These mice were sensitized using the strong contact sensitizer 2,4-dinitrochlorobenzene (DNCB), which resulted in a significant increase in ear swelling in Nrf2-deficient mice compared with WT mice, suggesting that Nrf2 mitigates inflammation in this model (Table 2, El Ali et al., 2013). Additionally, when mice were treated with low concentrations of DNCB, inflammation was solely observed in Nrf2-deficient mice, further supporting the idea that Nrf2 protects against contact dermatitis. The protective effect of Nrf2 was not limited to DNCB, as several other chemical sensitizers also increased lymphocyte proliferation in Nrf2-deficient mice compared with WT mice (El Ali et al., 2013).

However, activation of Nrf2 is not protective against every contact sensitizer. Topical application of 1-fluoro-2,4-dinitrobenzene (DNFB), an immunogenic hapten, was found to elicit an antioxidant response through the expression of Nrf2 downstream targets. However, in Nrf2-deficient mice, contact hypersensitivity was not developed due to a compromised epidermal innate immune response as demonstrated by a decrease in IL-1 α and keratinocyte-intrinsic factor, which is necessary for the development of immune memory (Ogawa et al., 2020b). Furthermore, clinical relevance of Nrf2 involvement in ACD was also assessed in this study through analysis of patients with various congenital disorders that resemble allergic contact dermatitis. Increased expression of Nrf2 and its target gene, small proline-rich protein 2, were found in the epidermis of patients with Netherton syndrome and peeling skin syndrome as compared with healthy controls. Nrf2 has also been shown to play a role in maintaining contact hypersensitivity response in aging mice. Aging is associated with diminished T helper (Th)1-mediated responses, including contact hypersensitivity. The decline in the contact

hypersensitivity in aged mice was more pronounced in Nrf2-deficient mice and less pronounced with treatment of the Nrf2 activator sulforaphane (Kim et al., 2008). Overall, these studies suggest that Nrf2 plays a contributing or causative role in the development of contact dermatitis in some instances.

Polymorphonuclear neutrophils (PMNs) are involved in hapten-induced inflammation during sensitization through the activation of DC (Weber et al., 2015). Nrf2 implication in PMN recruitment to DNCB-sensitized skin in WT and Nrf2-deficient mice was reported by Helou's group. They found that Nrf2 plays an important role in controlling PMN recruitment to the skin, which was demonstrated by an increase in lymphocyte antigen 6 locus, complex G/C (Ly6G/C) + cells in the skin of Nrf2-deficient mice after sensitization with DNCB (Helou et al., 2019). Additionally, antioxidant genes, such as HO-1, glutamate-cysteine ligase catalytic subunit (GCLC), and NQO1, were upregulated in WT mice compared with Nrf2-deficient mice sensitized with DNCB. Notably, they discovered an upregulation in the efferocytosis receptor CD36 on macrophages in the skin of sensitized WT mice, which was not observed in Nrf2-deficient mice. Overall, this study indicates that Nrf2 regulates neutrophil recruitment during the sensitization phase in this model of contact dermatitis.

A proteomics study was performed to assess potential biomarkers of activated DC by contact allergens, such as DNCB, cinnamaldehyde, and nickel (II) sulfate. Bone marrow-derived dendritic cells (BMDCs) from WT and Nrf2-deficient mice were used to determine the role of Nrf2. Several proteins, specifically stress response proteins such as glutamate-cysteine ligase modifier subunit (GCLM) and HO-1, were upregulated in WT cells treated with cinnamaldehyde and DNCB. Conversely, in Nrf2-deficient BMDCs, many of the proteins were not induced, suggesting Nrf2-dependent regulation (Mussotter et al., 2016). Taken together, the data from this study suggest an important role for Nrf2 in the induction of antioxidant genes by contact allergens in dendritic cells.

Current Challenges, Knowledge Gaps, and Future Directions. Based on studies dating back to the 2010s, there has been interest in developing Nrf2 activators for the treatment of allergic airway disease and skin allergy. However, the available therapeutic agents for this mechanism are limited in their practicality. Some of these therapeutic agents have related risks and are shown to have low efficacy and bioavailability (Egbujor et al., 2021). One of the challenges of these therapeutic agents is the lack of accurate and appropriate pharmacokinetic and pharmacodynamic reports, as well as safety profiles for administration.

Currently, there are several FDA-approved Nrf2 activators for clinical use. Dimethyl fumarate is approved for relapsing-remitting multiple sclerosis (Tecfidera) and internationally for psoriatic arthritis (Fumaderm). Recently, monomethyl fumarate (Bafiertam) and diroximel fumarate (Vumerity) have been approved for relapsing-remitting multiple sclerosis (Hoogendoorn et al., 2021). These drugs have also been tested in *in vitro* studies, which indicate that they induce Nrf2 target genes and would potentially have an antioxidant effect. Likewise, these drugs have been shown to ameliorate the symptoms of other chronic diseases, such as diabetes, chronic kidney disease, asthma, and dermatitis (Seidel and Roth, 2013; Hu et al., 2018; Zhao and Wen, 2018; Ogawa et al., 2020a).

One of the major challenges in Nrf2 drug development is the lack of specificity of Nrf2 activators reacting with protein thiol targets. Most Nrf2 activators that have been approved or are under clinical development aim to react with Cys151 of the N-terminal domain of Keap1. However, these Nrf2 activators have off-target effects and accompanied toxicity due to covalent and indiscriminate alkylation of thiol groups on other proteins (Gazaryan and Thomas, 2016; Robledinos-Anton et al., 2019). In addition, the use of many of these drugs is limited by short half-life (Hoogendoorn et al., 2021). Some synthetic drug candidates,

such as fumaric acid esters, sulforaphane, and nitro fatty acids have been considered as therapeutic agents and tested in clinical trials; however, they present inconsistent effects related to poor absorption, metabolism, and excretion (Egbujor et al., 2021). In addition to the covalent alkylation to thiol groups of Keap1, Nrf2 activation through a noncovalent inhibition of the Nrf2-Keap1 interaction is also being considered for therapeutic use (Satoh and Lipton, 2017).

Despite these challenges, there are ongoing clinical trials studying the pharmacokinetics of synthetic and naturally derived Nrf2 activators and inhibitors (Robledinos-Anton et al., 2019). Future work should be directed toward finding compounds or prodrugs with good pharmacokinetic/pharmacodynamic profiles that contain a more specific reaction to key thiol groups of Keap1 to avoid systemic side effects.

Authorship Contributions

Wrote or contributed to the writing of the manuscript: Rockwell, Jin, Boss, Kaiser, Awali.

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