Resveratrol Inhibits Paraquat-Induced Oxidative Stress and Fibrogenic Response by Activating the Nuclear Factor Erythroid 2-Related Factor 2 Pathway

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ABSTRACT
Nuclear factor erythroid 2-related factor 2 (Nrf2) is an antioxidant-activated transcription factor that recently emerged as a critical regulator of cellular defense against oxidative and inflammatory lesions. Resveratrol (Res) is a natural phytoalexin that exhibits multiple therapeutic potentials, including antioxidative and anti-inflammatory effects in animals. Paraquat (PQ) is the second most widely used herbicide worldwide, but it selectively accumulates in human lungs to cause oxidative injury and fibrosis with high mortality. Here, we analyzed the molecular mechanism of the fibrogenic response to PQ and its inhibition by Res and Nrf2. PQ dose-dependently caused toxicity in normal human bronchial epithelial cells (BEAS-2B), resulting in mitochondrial damage, oxidative stress, and cell death. Res at 10 μM markedly inhibited PQ toxicity. PQ at 10 μM stimulated production of inflammatory and profibrogenic factors (tumor necrosis factor α, interleukin 6, and transforming growth factor β1) and induced the transformation of normal human lung fibroblasts (WI38-VA13) to myofibroblasts; both effects were inhibited by Res. Res strongly activated the Nrf2 signaling pathway and induced antioxidant response element-dependent cytoprotective genes. On the other hand, knockout or knockdown of Nrf2 markedly increased PQ-induced cytotoxicity, cytokine production, and myofibroblast transformation and abolished protection by Res. The findings demonstrate that Res attenuates PQ-induced reactive oxygen species production, inflammation, and fibrotic reactions by activating Nrf2 signaling. The study reveals a new pathway for molecular intervention against pulmonary oxidative injury and fibrosis.

Introduction
Pulmonary fibrosis is an irreversible stage of a large heterogeneous group of chronic lung diseases with a 5-year mortality rate larger than 50% (Husain and Kuman, 2005). Fibrosis in the lungs can result from genetic, infectious, autoimmune, idiopathic, and cancer pathology or arise from exposure to environmental and occupational agents, such as fibers and particles, metals, pesticides, and anticancer drugs (Bus and Gibson, 1984; Castranova and Vallyathan, 2000; Husain and Kuman, 2005; Rom, 2007). The underlying mechanism for lung fibrosis is unclear for the most part. Several cellular responses are commonly observed during lung fibrosis, including injury to airway and alveolar epithelia, macrophage activation, and transformation of fibroblasts into myofibroblasts. Myofibroblasts synthesize collagen 1 and α smooth muscle actin (αSMA) to promote fibrosis and scar formation (Fichtner-Feigl et al., 2006; Wynn, 2008; Bonner, 2010). The molecular events governing these fibrogenic reactions remain poorly understood. Although research trials are ongoing, there is no evidence that any medication can significantly help lung fibrosis at the present time.

Paraquat (PQ) is a highly effective, fast-acting, and nonselective herbicide widely used in the world. Human exposure to PQ by either respiratory or systemic route leads to the
accumulation of PQ in the lungs, resulting in pulmonary edema, bronchial and alveolar destruction, and ultimately fibrosis with high mortality, which is in part caused by the lack of a specific antidote (Bus and Gibson, 1984). Chronic exposure to PQ is associated with liver damage, kidney failure, and Parkinsonian lesions in addition to fibrosis (Ossowska et al., 2006; Tanner et al., 2011). Upon entering cells, PQ undergoes cyclic single-electron reduction/oxidation through its quaternary ammonium nitrogen atoms and bipyridyl ring, producing reactive oxygen species (ROS) and PQ radicals. Redox cycling is believed to play an important role in initiating lung damage and fibrosis by PQ. How the oxidative signals from PQ interact with the pathways that underlie lung fibrogenic response is poorly understood.

Resveratrol (Res) is a phytoalexin polyphenol produced naturally by several plants when under attack by bacterial and fungal pathogens. Res exhibits multiple health-promoting properties including antioxidative, anticancer, anti-inflammatory, antiaging, blood sugar-lowering, and beneficial cardiovascular effects (Jang et al., 1997; Duffy and Vita, 2003; Leonard et al., 2003; Su et al., 2006; Valenzano et al., 2006; Elmali et al., 2007). Res alleviated bleomycin-induced lung injury in rats that typically progresses to fibrosis otherwise (Sener et al., 2007). Although the mechanism of protection by Res against bleomycin lung toxicity remains unclear, its beneficial effects on the vascular endothelium and lung epithelium involved activation of the nuclear factor erythroid 2-related factor 2 (Nrf2) pathway (Kode et al., 2008; Ungvari et al., 2010).

Nrf2 is a cap “n” collar basic leucine zipper transcription factor ubiquitously expressed in mammalian cells. Nrf2 controls the antioxidant response element (ARE)-mediated expression of cellular detoxification enzymes and antioxidant proteins in response to a wide range of oxidant/antioxidant/electrophilic stimuli to protect the body (Leung et al., 2003; Talalay et al., 2003; Kobayashi et al., 2004; Kensler et al., 2005).

Fig. 1. Inhibition of PQ toxicity by Res. A, cytotoxicity. BEAS-2B cells were treated with PQ or PQ + Res (10 μM) for 24 h in triplicate. Cytotoxicity was detected by using CellTiter 96 AQueuos One Solution Reagent. Data represent means from three samples. Inset, apoptosis (caspase 3/7 activation) was measured in cells treated with PQ at 10 μM for 24 h. Caspase activity was expressed in RLU. B, ROS production. Cells were treated with Res (10 μM) for 16 h then with or without PQ (20 μM) for 24 h. ROS production was measured by using fluorophore DHE and expressed in RFU. Data represent means ± S.D. from three samples. C, mitochondrial damage. Cells were treated as in B. Mitochondrial damage was assayed by using fluorophores TMRE (green) and MTDR (red). DAPI (blue) was used to counterstain the nucleus. Composite images show the mitochondria in yellow (merge of green and red). Bar size, 20 nm. D, quantification of TMRE and MTDR fluorescence. *, p < 0.05; **, p < 0.01.
Nrf2 and its binding protein Keap1 are redox sensors. Modification of critical cysteine residues in Keap1 and Nrf2 by oxidants and electrophiles leads to suppression of the ubiquitination and proteasomal degradation of Nrf2 (Kobayashi et al., 2004; He et al., 2006; He and Ma, 2009, 2010; Itoh et al., 2010; Taguchi et al., 2011). Activated Nrf2 translocates into the nucleus, dimerizes with small Maf proteins, and binds to ARE to up-regulate the transcription of cytoprotective genes. Nrf2-deficient mice have a tendency to develop autoimmune lesions and sponge-form leukoencephalopathy (Ma et al., 2006; Hubbs et al., 2007) and have increased susceptibility to toxicant-induced diseases, such as benzo[a]pyrene-induced cancer (Ramos-Gomez et al., 2001), metal toxicity (He et al., 2007, 2008), premature ovarian failure by 4-vinylcyclohexene diepoxide (Hu et al., 2006), and diabetic cardiomyopathy induced by high glucose (He et al., 2009). It is noteworthy that Nrf2 null mice are vulnerable to bleomycin and hypoxia-induced airway inflammation and lung fibrosis (Cho et al., 2004).

The purpose of the study was 3-fold: 1) to elucidate the molecular mechanism by which PQ damages human lung cells and induces fibrogenic response; 2) to examine the effect of Res on PQ pulmonary toxicity; and 3) to analyze the role and mechanism of action of Nrf2 in these processes. We found that PQ induced mitochondrial damage that, together with PQ redox cycling, caused heightened oxidative stress, leading to lung epithelial cell death, production of profibrogenic cytokines and growth factors, and myofibroblast transformation. Res effectively prevented PQ toxicity and blocked the fibrogenic reactions. Finally, Nrf2 was found to be critical for defense against PQ toxicity in normal cells and the protective effects of Res. The findings revealed a molecular pathway for

![Graph A](image1.png)

**Graph A**: TNFα enhancer

Reported Luminescence (RLU)

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![Graph B](image2.png)

**Graph B**: TNFα (pg/ml)

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![Graph C](image3.png)

**Graph C**: IL-6 (pg/ml)

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![Graph D](image4.png)

**Graph D**: Relative Amount of TGFβ1 mRNA

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**Fig. 2.** Suppression of PQ-induced expression of TNFα, IL-6, and TGFβ1 by Res. A. TNFα promoter/luciferase reporter expression. Macrophages stably transfected with the TNFα promoter/luciferase reporter plasmid were treated with LPS (1 μg/ml, positive control), PQ (10 μM), Res (10 μM), LPS + Res, or PQ + Res for 16 h. Luciferase reporter expression was measured and expressed in RLU. B and C, secretion of TNFα (B) and IL-6 (C). Macrophages were pretreated with Res (10 μM) for 16 h followed by PQ (10 μM) for another 24 h (Res + PQ). LPS (1 μg/ml, 5 h) was a positive control for induction. The culture medium was assayed for TNFα (B) or IL-6 (C) proteins by flow cytometry using TNFα and IL-6 Flex beads. D, expression of TGFβ1 mRNA. WI38-VA13 cells were treated with PQ (10 μM) for 24 h or Res (10 μM) for 16 h followed by PQ for another 24 h (Res + PQ). Total RNA was assayed for TGFβ1 mRNA expression by using real-time PCR. Actin was used as a loading control. Data represent means ± S.D. from three samples. *p < 0.05; **p < 0.1; ***p < 0.001.
intervention against PQ lung fibrosis and, potentially, other fibrotic diseases.

Materials and Methods

Reagents and Cell Culture. Reagents and cells were purchased from commercial vendors: PQ (N,N'-dimethyl-4,4'-bipyridinium dichloride; Thermo Fisher Scientific, Waltham, MA); Res (3,5,4'-trihydroxy-trans-stilbene; Enzo Life Sciences, Inc., Farmingdale, NY); tetramethylrhodamine ethyl ester (TMRE) and lipopolysaccharide (LPS) (Sigma, St. Louis, MO); MitoTracker deep red 633 (MTDR) and cell culture media (Invitrogen, Carlsbad, CA); and BEAS-2B, WI38-VA13, and RAW264.7 cells (American Type Culture Collection, Manassas, VA). Mouse embryonic fibroblast (MEF) cells were derived from Nrf2 wild-type (WT) and knockout (KO) mice and immortalized as described elsewhere (Bi et al., 2004; He and Ma, 2009). MEFs were cultured in Dulbecco’s modified Eagle’s medium with 10% FBS and 5% CO2. BEAS-2B cells were cultured in basal medium Eagle’s medium with additives from Lonza Walkersville, Inc. (Walkersville, MD). WI38-VA13 cells were cultured in α-modified Eagle’s medium with 10% FBS. Raw264.7 cells were grown in Dulbecco’s modified Eagle’s medium with 10% FBS.

Cytotoxicity Assay. The CellTiter 96 AQeone, One Solution Cell Proliferation Assay kit (Promega, Madison, WI) was used to detect cell proliferation/cytotoxicity following the manufacturer’s instructions. Cells (1 × 10^5) were seeded in 96-well plates overnight. Absorbance at 490 nm (A490nm) was recorded, and means and S.D. were calculated from three samples for each treatment.

ROS Detection. Cells in an eight-well chamber slide were added with 5 μM dihydroethidium (DHE; fluorophore for ROS; Invitrogen) 30 min before the end of treatment. Cells were fixed with 4% paraformaldehyde and mounted with a mounting solution containing DAPI that stains the nucleus. Images were taken by using a Zeiss LSM510 confocal microscope (Carl Zeiss Inc., Thornwood, NY) with a rhodamin-DAPI setting and fixed exposure time. Fluorescence intensity was quantified by using Optimus version 6.51 software (Media Cybernetics Inc., Bethesda, MD) and expressed in relative fluorescence units (RFU). Means and S.D. were calculated from five separate fields for each treatment.

Measurement of Mitochondrial Inner Membrane Potential. Cells cultured in an eight-well chamber slide were added with 50 nM TMRE and 1 μM MTDR 45 min before the end of the treatment. Cells were fixed with 4% paraformaldehyde and mounted with a mounting solution containing DAPI for fluorescence microscopy. Quantification of fluorescence intensity was performed as described above for ROS detection. Alternatively, TMRE and MTDR fluorescence was measured by flow cytometry, and data were expressed in RFU.

Immunoblotting. Cells were lysed on ice with radioimmunoprecipitation assay buffer containing protease and phosphatase inhibitors for 30 min. Cell lysate (40 μg) was immunoblotted with anti-Nrf2 or actin antibodies (Santa Cruz Biotechnology, Inc., Santa Cruz, CA). Protein bands were visualized by using enhanced chemiluminescence detection reagents from GE Healthcare (Chalfont St. Giles, Buckinghamshire, UK).

Detection of Cytokine. Cells were treated as shown in figure legends, and cell culture medium was collected. TNFα and IL-6
cytokines were detected by flow cytometry using mouse Flex TNFs and IL-6 beads (BD Biosciences, San Diego, CA) following the manufacturer’s instructions.

**Immunofluorescent Staining.** Cells in eight-well chamber slides were fixed with 4% paraformaldehyde, permeated with 0.5% Triton X-100, and stained with anti-αSMA antibody (Sigma) overnight at 4°C, followed by Alexa Fluor 594-conjugated second antibody (Invitrogen) for 1 h, avoiding light. Fluorescence images were taken and quantified as described above for ROS detection.

**Apoptosis.** BEAS-2B cells cultured in 96-well plates were added with 100 μl of the Caspase-3-Glo 3/7 reagent (Promega) and cultured at 37°C for 1 h. Luminescence was measured in a plate-reading luminometer and expressed in relative light units (RLU).

**Reporter Assay.** RAW264.7 cells stably transfected with a TNFα promoter/luciferase reporter construct were as described previously (Ma et al., 2003). Cells were treated and lysed with a passive reporter lysis buffer (Promega). Cell lysate was centrifuged briefly to remove cell debris. Twenty microliters of supernatant was mixed with 100 μl of luciferase reagent (Promega), and luciferase activity was detected by using a TD 20/20 luminometer ( Molecular Devices, Sunnyvale, CA). Luciferase activity was in RLU and normalized with protein concentrations.

**Real-Time PCR.** Cells were treated as shown in figure legends. Total RNA was isolated by using a Qiagen RNA mini kit (Qiagen, Valencia, CA). Five micrograms of total RNA was reverse-transcribed into cDNA with reverse transcriptase III (Invitrogen). TOFp1 mRNA was amplified with gene-specific primers (primer sequences available on request) by using real-time PCR as described previously (He et al., 2009). Actin was used as an internal control.

**siRNA Transfection.** Cells were transfected with either the control siRNA (Santa Cruz Biotechnology, Inc) or Nrf2 siRNA specific for mouse Nrf2 (Invitrogen) as described previously (He et al., 2009). Knockdown of Nrf2 was confirmed by immunoblotting of Nrf2.

**Northern Blotting.** Total RNA of 3 μg each was fractionated in a 1.2% formaldehyde agarose gel, transferred to a supercharged nylon membrane, and blotted with the digoxigenin-labeled riboprobe preparations with digoxigenin-labeling reagents (Roche Applied Science, Indianapolis, IN). Plasmid constructs for riboprobes of Nqo1, Ho1, and actin were verified by sequencing. Northern signals were visualized by chemiluminescence using a digoxigenin RNA detection kit with CDP Star as a substrate (Roche Applied Science).

### Results

**Res Inhibits PQ-Induced Cell Death, ROS Production, and Mitochondrial Damage in Human Lung Cells.** Exposure to PQ in the lungs causes extensive bronchial and alveolar destruction before fibrosis. BEAS-2B cells, a normal human bronchial epithelium cell line, were used to examine PQ cytotoxicity and the effect of Res. PQ at concentrations of 10^{-5} to 10^{-3} M induced cell toxicity dose-dependently (Fig. 1A). Toxicity was largely attributed to apoptosis as revealed by caspase 3/7 activation (Fig. 1A, inset). Res at 10 μM significantly reduced PQ toxicity at concentrations of 10^{-5} and 10^{-4} M.

PQ undergoes redox cycling producing O_2^− and other radicals to initiate cell killing in plants and animals. Incubation of BEAS-2B cells with PQ (20 μM, 24 h) dramatically increased ROS production (5-fold) detected by DHE, a fluorophore specific for O_2^− (Fig. 1B). Res (Res + PQ; pretreated at 10 μM before PQ) significantly reduced the amount of PQ-induced ROS (~70%).

The mitochondria consume approximately 90% of the oxygen in cells for ATP synthesis and are a major source of ROS under both normal and pathological conditions. The observation that PQ substantially increased ROS production, whereas Res largely blocked PQ-induced ROS production suggests the involvement of mitochondria in PQ and Res effects. TMRE and MTDR are cationic lipophilic fluorophores that selectively accumulate in mitochondrial matrix in a mitochondrial inner membrane potential-dependent manner. Strong TMRE and MTDR fluorescence signals were found in the cytoplasm of control cells, indicating accumulation of the dyes in the mitochondrial matrix; however, the fluorescence was largely reduced and became diffusive upon treatment with PQ (20 μM; 24 h), indicating mitochondrial damage (Fig. 1, C and D). On the other hand, pretreatment with Res for 16 h followed by PQ restored the fluorescence intensity, indicating PQ-induced mitochondrial damage was largely prevented by Res.

**Res Blocks PQ-Induced Fibroblast-to-Myofibroblast Transformation by Suppressing the Production of Inflammatory and Profibrogenic Factors.** Fibrogenic agents induce the production and secretion of inflammatory and profibrogenic cytokines and growth factors from macrophages, epithelial cells, and fibroblasts. The soluble factors function as paracrines and autocrines to stimulate inflammation and fibrosis. To examine the effects of PQ and Res on the expression of profibrogenic factors, a luciferase reporter under the control of mouse TNFα promoter was examined for induction in RAW264.7 cells, a widely used macrophage cell line (Fig. 2A). PQ (10 μM; 16 h) induced reporter expression.

![Fig. 4. Activation of Nrf2. A, Nrf2 protein stabilization. BEAS-2B cells were treated with Res (10 μM), PQ (10 μM), or PQ + Res for 24 h. Total cell lysates were immunoblotted with an anti-Nrf2 antibody. Actin was used as loading control. B, induction of Nqo1 and Ho1 mRNA. BEAS-2B cells were treated with tBHQ (30 μM), Res (10 μM), PQ (10 and 100 μM), Res + PQ (10 μM), or Res + PQ (100 μM) for 24 h. Total RNA was fractionated on 1.2% formaldehyde agarose gel. RNA bands were blotted with digoxigenin-labeled probes for Nqo1, Ho1, and actin and visualized with the CDP Star reagents and chemiluminescence.](https://jpet.aspetjournals.org/doi/abs/10.1124/jpet.117.233518)
Res itself (10 μM; 16 h) did not affect reporter expression, but cotreatment with Res blocked reporter induction by PQ. Res also weakly, but significantly, reduced reporter induction by LPS, a known inducer of TNFα. Expression of endogenous genes was examined. PQ (10 μM; 24 h) induced production of TNFα (Fig. 2B) and IL-6 (Fig. 2C) proteins in macrophages and TGFβ1 mRNA in fibroblasts (Fig. 2D). Pretreatment with Res at 10 μM for 16 h totally blocked induction of TNFα and IL-6 proteins and significantly diminished the induction of TGFβ1 mRNA by PQ.

Transformation of fibroblast to myofibroblast is a critical cellular event in fibrosis because the latter synthesizes collagen I and αSMA, two major proteins involved in fibrosis and scar formation. WI38-VA13 cells, a normal human lung fibroblast cell line, were used to assess the effects of PQ and Res on myofibroblast transformation. PQ (10 μM; 24 h) induced a significant increase (>3-fold) in the amount of αSMA, indicating transformation of fibroblasts to myofibroblasts (Fig. 3). Res itself (10 μM) did not affect αSMA expression, but pretreatment with Res markedly reduced PQ-induced increase in αSMA. Thus, PQ stimulated myofibroblast transformation, whereas Res blocked the transformation.

**Res Activates Nrf2 to Induce Cytoprotective Genes.**
Nrf2 controls the cellular defense against oxidative and electrophilic chemicals and lesions. The profound oxidative effect by PQ and its inhibition by Res in lung cells raised the possibility that Nrf2 is involved in the actions of PQ and Res. As shown in Fig. 4A, Res (10 μM; 24 h) strongly and PQ (10 μM; 24 h) weakly increased the protein level of Nrf2 in BEAS-2B cells, indicating activation of Nrf2. Cotreatment with Res and PQ further increased Nrf2 protein, suggesting a synergy between Res and PQ for Nrf2 activation. Northern blotting was performed to analyze the induction of Nqo1 and Hsp70, two representative target genes of Nrf2 (Fig. 4B). tBHQ (30 μM), a known inducer of Nqo1 and Hsp70, strongly induced the genes. Res (10 μM) induced Nqo1 similarly to tBHQ and Hsp70 to a lesser extent. PQ induced both genes at 100 but not 10 μM. Therefore, Res and, to a lesser extent, PQ activated Nrf2 signal transduction to induce cytoprotective genes.

**Nrf2 Controls Defense against the Cytotoxicity, Oxidative Stress, and Mitochondrial Damage of PQ and Mediates Protection by Res.**
The role of Nrf2 in PQ toxicity and protection by Res was examined by comparing Nrf2 WT and KO MEFs. PQ at 10⁻⁷ and 10⁻⁴ M caused significant toxicity in WT cells, and Res at 10 μM significantly reduced the toxicity (Fig. 5A) in agreement with the data for BEAS-2B cells (Fig. 1A). In KO cells, PQ induced significantly a higher level of toxicity than wild type and Res failed to rescue. The results indicate that Nrf2 is required for defense against PQ toxicity in normal cells and for protection by Res.

In WT cells, PQ (10 μM) induced a dramatic increase of ROS production. Res (10 μM) alone did not significantly change the ROS level, but Res cotreatment blocked PQ-induced ROS production (Fig. 5, B and C). In KO cells, the basal ROS production was increased (compare control KO with control WT); PQ induced a high level of ROS; Res alone did not significantly change ROS level; and Res failed to block PQ-induced ROS production. The data revealed that Nrf2 is critically involved in the control of ROS production in cells exposed to PQ and is required for the suppression of PQ-induced oxidative stress by Res.

![Fig. 5](image-url) Nrf2-dependent inhibition of PQ toxicity. A, cell toxicity. Nrf2 WT and Nrf2 KO MEF cells were treated with PQ or PQ + Res (10 μM) for 24 h in triplicate. Cytotoxicity was measured and analyzed as described for Fig. 1. B, ROS production. Nrf2 WT and Nrf2 KO MEF cells were treated with PQ (10 μM), Res (10 μM), or PQ + Res for 24 h. ROS fluorescent probe DHE was added 30 min before the end of treatment to detect ROS. DAPI was used to counterstain the nucleus. Bars, 20 nm. C, quantification of fluorescence intensity of B. D and E, BEAS-2B cells were transfected with control siRNA or Nrf2 siRNA. The cells were then treated as described for B. D, knockdown of Nrf2 was examined by immunoblotting of Nrf2 with actin as the loading control. E, mitochondrial damage was assessed by using TMRE and flow cytometry. Data represent means ± S.D. from three samples.* p < 0.05; **, p < 0.01.
The effect of Nrf2 on PQ toxicity to BEAS-2B mitochondria was examined by using siRNA knockdown of Nrf2 (Fig. 5, D and E). Knockdown of Nrf2 significantly reduced the protein level of Nrf2 (Fig. 5D). Knockdown of Nrf2 or treatment with Res (10 µM) alone did not alter the mitochondrial inner membrane integrity, but PQ (10 µM) significantly reduced TMRE accumulation in mitochondrial matrix in both control and Nrf2 knockdown cells (Fig. 5E). Cotreatment with Res rescued the cells from PQ-induced mitochondrial damage in WT but not Nrf2 knockdown cells, indicating Nrf2 is required for Res protection against PQ toxicity to mitochondria.

**Nrf2 Inhibits Induction of Profibrogenic Factors and Transformation of Myofibroblast by PQ.** The effect of Nrf2 on the regulation of inflammatory and profibrogenic cytokine and growth factor expression by PQ and Res was examined in macrophages transfected with control or Nrf2 siRNA (Fig. 6). Nrf2 protein was significantly diminished in macrophages upon knockdown of Nrf2 (Fig. 6A). PQ (10 µM; 24 h) induced TNFα secretion from macrophages, and induction was significantly higher in Nrf2 knockdown cells, suggesting Nrf2 suppresses TNFα expression in macrophages (Fig. 6B). Res (pretreated for 16 h at 10 µM) blocked TNFα induction by PQ in control but not Nrf2 knockdown cells. Similar effects of Nrf2 knockdown were observed for the induction of IL-6 protein (Fig. 6C) and TGFβ1 mRNA (Fig. 6D) by PQ and inhibition by Res. Thus, Nrf2 inhibited PQ induction of the cytokines and growth factors in WT cells, and Nrf2 is required for Res to block induction by PQ.

PQ (10 µM; 24 h) induced transformation of Nrf2 WT fibroblasts into myofibroblasts as evidenced by marked increase in the synthesis of αSMA (Fig. 7). Res (10 µM) alone did not affect αSMA expression, but pretreatment with Res at 10 µM for 16 h markedly inhibited induction of αSMA by PQ. In Nrf2 KO fibroblasts, PQ induced significantly higher amounts of αSMA than in WT cells, and this increase was not affected by Res pretreatment. Therefore, Nrf2 inhibited PQ-induced myofibroblast transformation, and Nrf2 was required for suppression of PQ-induced fibrotic reaction by Res.

![Figure 6](https://example.com/fig6.png)

**Fig. 6.** Effect of Nrf2 on cytokine expression. RAW264.7 macrophages were transfected with control siRNA or Nrf2 siRNA and then treated with PQ (10 µM) or Res + PQ (pretreated with Res at 10 µM for 16 h followed by PQ for 24 h). A, knockdown of Nrf2 is shown by immunoblotting. B and C, TNFα (B) and IL-6 (C) proteins in the culture medium were detected by flow cytometry. D, induction of TGFβ1 mRNA was examined by real-time PCR. *p < 0.05; **p < 0.01; ***p < 0.001.
Discussion

The hallmark of PQ pulmonary toxicity is extensive tissue damage followed by fibrosis after either respiratory or systemic exposure (Bus and Gibson, 1984), which resembles the toxicity of a number of pulmonary toxicants, such as bleomycin and nitrofurantoin. In the case of PQ, the quaternary ammonium bipyridyl structure makes it prone to cyclic one-electron reduction/oxidation that causes production of ROS and depletion of reducing equivalents NADPH and GSH, both contributing to oxidative stress in cells. Thus, PQ serves as a valuable model compound for studying pulmonary toxicity and fibrosis, in particular, oxidative stress in lung fibrosis.

In this study, we analyzed the mechanism of the fibrogenic response to PQ and its inhibition by Res and Nrf2 in lung cells. Taken together, the findings illustrate a molecular model in which PQ induces oxidative stress and fibrotic reactions in lung cells, whereas Res suppresses PQ-induced oxidative stress and fibrogenic responses by activating the Nrf2 pathway (Fig. 8). PQ induces oxidative stress through both redox cycling and mitochondrial damage. Oxidative stress stimulates fibrogenic responses by promoting production and secretion of inflammatory and profibrogenic factors from macrophages and other lung cells, transformation of fibroblasts to myofibroblasts, and synthesis of αSMA and collagen I. Res activates Nrf2 and induces cytoprotective genes to inhibit PQ-induced ROS production, mitochondrial damage, cytokine and growth factor expression, and myofibroblast transformation.

The mitochondria are a major organelle for oxygen consumption and ROS production in cells. We found that mitochondria are a major target of PQ and damage to mitochondria contributes to oxidative stress induced by PQ. In this regard, PQ-induced mitochondrial damage and PQ redox cycling form a vicious cycle to cause heightened oxidative stress in lung cells. The observations that loss of Nrf2 in cells exacerbated PQ-induced ROS production and mitochondrial toxicity, whereas antioxidant Res blocked PQ oxidative stress and toxicity, support the notion that oxidative stress and mitochondrial damage are major mechanisms by which PQ causes lung damage and fibrosis.

Res is one of the promising natural products with multiple health-promoting effects (Vang et al., 2011). In this study, Res was shown, for the first time, to inhibit PQ-induced oxidative damage, expression of profibrogenic cytokines and growth factors, and transformation of fibroblasts into myofibroblasts effectively. Several lines of evidence support that Nrf2 mediates the protective effect of Res against PQ lung
toxicity and fibrosis. First, Res elevated the Nrf2 protein level and induced Nrf2/ARE-controlled cytoprotective genes Nqo1 and Ho1 in lung cells. Second, PQ induced oxidative stress and fibrogenic reactions that were more severe in Nrf2 KO than WT cells. Third, Res could reverse or attenuate PQ-induced ROS production and fibrogenic responses in WT but not KO cells. Finally, a body of literature exists indicating a critical role of Nrf2 in Res functions. For instance, activation of Nrf2 is an important mechanism by which Res exerts its protective effects in the vascular endothelium (Ungvari et al., 2010). Res protects against cigarette smoke-mediated oxidative damage by activating Nrf2 in human lung epithelial cells (Kode et al., 2008), and both Nrf2 and Res protect against bleomycin-induced lung damage and fibrosis (Cho et al., 2004; Sener et al., 2007).

Cytokines and growth factors are involved in both the early inflammatory and late fibrotic phases of lung fibrosis through paracrine and autocrine mechanisms. TNFα is one of the early cytokines consistently found in animal models of pulmonary fibrosis and plays a cardinal role in the pathogenesis of this disease. TNFα activates nuclear factor-κB and activator protein 1 that regulate the proliferation of fibroblasts and expression of cellular matrix deposition genes, such as matrix metalloproteinase-9. IL-6 promotes fibrogenesis either alone or in concert with TNFα. IL-6 mRNA level is elevated in idiopathic pulmonary fibrosis (Smith et al., 1998). Overexpression of IL-6 in the lungs does not seem to be sufficient to induce marked fibrosis, but an abnormal IL-6 level has been consistently detected in animal and human lung fibrosis samples (Baecher-Allan and Barth, 1993; Moodley et al., 2003). We found that PQ significantly increased the expression of TNFs and IL-6 in macrophages, but pretreatment with Res prevented induction of TNFα and IL-6 by PQ.

TGFβ1 is a critical regulator of cell growth and differentiation as well as matrix protein production. Elevated production of TGFβ1 has been associated with human and rodent chronic inflammatory and fibrotic diseases. For instance, bleomycin-induced pulmonary fibrosis is accompanied by enhanced expression of TGFβ1, collagen 1, and αSMA (Zhang et al., 1996). TGFβ induces the proliferation and differentiation of ex vivo human lung fibroblasts into myofibroblasts through extracellular signal-regulated kinase/Akt inhibition and Pten restoration, whereas Res inhibits this process (Fagone et al., 2011). PQ increased TGFβ1 production, but pretreatment with Res for 16 h could prevent PQ-induced TGFβ1 expression.

The primary effector cells in fibrosis are myofibroblasts, which are highly synthetic for collagen 1, have a contractile phenotype, and are characterized by de novo synthesis of αSMA stress fiber. Myofibroblasts may derive from activation/proliferation of resident lung fibroblasts, through epithelial-mesenchymal differentiation, or via recruitment of circulating fibroblastic stem cells. Resting fibroblasts express only β- and γ-actin isoforms (Herman, 1993; Khahtila, 2001), whereas activated fibroblasts synthesize αSMA. Res could inhibit αSMA expression in hepatic stellate cells (Kawada et al., 1998). Olson et al. (2005) demonstrated an inhibitory role of Res in cardiac fibroblast proliferation and myofibroblast formation by reducing αSMA expression and stress fiber organization in cardiac fibroblasts. Res also alleviates bleomycin-induced pulmonary fibrosis (Sener et al., 2007). We found that PQ treatment for 24 h elevated αSMA expression.
in lung fibroblast WI38VA13 cells, but pretreatment with Res for 16 h inhibited PQ-induced αSMA expression, providing the first evidence that Res could effectively prevent PQ-induced fibrogenic response in lung cells. These data provide a rationale for exploring the therapeutic and chemoprotective potentials of Res for the treatment and prevention of pathologic conditions involving pulmonary fibrosis.

Authorship Contributions

**Participated in research design:** He, Wang, Sklarz, Bi, and Ma. **Conducted experiments:** He, Wang, and Ma. **Contributed new reagents or analytic tools:** Sklarz and Bi. **Performed data analysis:** He, Wang, and Ma. **Wrote or contributed to the writing of the manuscript:** He, Wang, and Ma.

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