

The Role of NADPH Oxidase 1–Derived Reactive Oxygen Species in Paraquat-Mediated Dopaminergic Cell Death

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Abstract

Oxidative stress is the common downstream effect of a variety of environmental neurotoxins that are strongly implicated in the pathogenesis of Parkinson's disease. We demonstrate here that the activation of NADPH oxidase 1 (Nox1), a specialized superoxide-generating enzyme complex, plays a key role in the oxidative stress and subsequent dopaminergic cell death elicited by paraquat. Paraquat increased the expression of Nox1 in a concentration-dependent manner in rat dopaminergic N27 cells. Rac1, a key component necessary for Nox1-mediated superoxide generation, also was activated by paraquat. Paraquat-induced reactive oxygen species generation and dopaminergic cell death were significantly reduced after pretreatment with apocynin, a putative NADPH oxidase inhibitor, and Nox1 knockdown with siRNA. Male C57BL/6 mice received intraperitoneal (IP) injections of paraquat (10 mg/kg) once every 3 days and showed increased Nox1 levels in the substantia nigra as well as a 35% reduction in tyrosine hydroxylase–positive dopaminergic neurons 5 days after the last injection. Preadministration of apocynin (200 mg/kg, IP) led to a significant decrease in dopaminergic neuronal loss. Our results suggest that Nox1-generated superoxide is implicated in the oxidative stress elicited by paraquat in DA cells, and it can serve as a novel target for pharmacologic intervention. *Antioxid. Redox Signal.* 11, 2105–2118.

Introduction

ALTHOUGH the specific etiology of Parkinson's disease (PD) remains largely elusive, aging, genetic susceptibility, and exposure to environmental toxic compounds contribute to the development of PD. Based on epidemiologic studies showing that persons living in rural areas, who farm and drink well water, have a higher PD incidence, it has been widely postulated that agricultural agents may be linked to PD pathogenesis (5, 10). Paraquat (1,1'-dimethyl-4,4'-bipyridinium dichloride, PQ), a widely used herbicide, is in this category and is considered a key risk factor. Epidemiologic studies demonstrated the association between prolonged exposure to PQ and increased risk for developing PD (28, 39). Although the molecular mechanisms governing PQ toxicity to the nigrostriatal dopaminergic (DA) system are still under extensive investigation, its abilities to produce superoxide *via* the redox cycling and to induce ROS generation by mitochondrial inhibition suggest PQ as an oxidative stressor (22). Various cellular reductases can catalyze the one-electron reduction of PQ to a cation radical, which, by transferring its

electron onto molecular oxygen, would readily form superoxide anion. The production of superoxide radical also regenerates a PQ parent compound (11, 22). This redox cycling has the potential to yield large amounts of reactive oxygen species (ROS) from relatively small concentrations of an agent (17, 22).

A growing body of evidence has demonstrated that oxidative stress is a key player in the pathogenesis of PD. DA neurons are extremely sensitive to insult. This selective vulnerability of the nigrostriatal system to oxidative stress is based on the following observations: (a) the generation of ROS during the oxidative metabolism of dopamine (25); (b) the capability of its quinone metabolites to adduct proteins containing a sulfhydryl group such as glutathione (6, 34); and (c) the high level of iron in the substantia nigra (SN) and globus pallidus that contributes to the highly reactive hydroxyl radical (OH[•]) generation *via* the Fenton reaction (21, 26). The idea is supported by studies of postmortem brain tissues from PD patients demonstrating high oxidative stress levels in the SN, marked by increased lipid peroxidation (20), protein (2) and DNA (60) oxidative damage, and decreased

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glutathione levels (46, 54). Mitochondrial dysfunction also has been implicated in PD pathogenesis, partly through an increase in ROS generation. Mitochondrial respiratory chain complex I activity is selectively decreased in the SN of PD postmortem brains, whereas other electron-transport complexes remain unchanged (51). Complex I inhibitors, such as 1-methyl-4-phenylpyridinium (MPP⁺), the reactive product of 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) and rotenone, cause oxidative stress followed by DA neuronal loss that replicates the hallmarks of PD in rodents (7, 53). Inflammation mainly caused by microglial activation in the central nervous system (CNS) is also responsible for the PD pathogenesis. Superoxide and nitrogen oxide (NO) are directly secreted from activated microglia and lead to DA neuronal degeneration (24, 30).

Nicotinamide adenine dinucleotide phosphate (NADPH) oxidase was originally discovered in phagocytes (50). In these cells, the enzyme is responsible for killing bacteria through the release of substantial quantities of superoxide into the phagosomes. Although several organelles including mitochondria and biological reactions physiologically generate ROS as a by-product, the NADPH oxidase (Nox) system functions primarily as a superoxide generator. The recent discovery of several isoforms of the Nox family reveals that this specialized ROS-generation system is not limited to phagocytes (14, 18, 55). Within the CNS, studies have focused primarily on microglial gp91^{phox} (Nox2). Nox2 knockout led to reduced DA toxicity caused by PQ in both *in vitro* and *in vivo* studies (47, 59). Recent studies have shown, however, that various Nox homologues are expressed in neurons as well as in astrocytes (33).

Based on this background, we investigated whether DA cells are equipped with the NADPH oxidase system, and if this system is readily activated by toxic insults such as PQ to produce ROS, and results in DA neurodegeneration.

Materials and Methods

Materials

Fetal bovine serum (FBS), horse serum, RPMI 1640, L-glutamine, trypsin/EDTA, and penicillin-streptomycin were purchased from GibcoBRL (Gaithersburg, MD). Phenylmethylsulfonyl fluoride (PMSF), Nonidet P-40 (NP-40), SP600125, Brij35, and bupropion were purchased from Sigma Chemicals (St. Louis, MO). Rabbit polyclonal anti-tyrosine hydroxylase (TH) was obtained from Protos Biotech (New York, NY), rabbit anti-Nox1 from Santa Cruz Biotechnology (Santa Cruz, CA), and rat anti-CD11b from Serotec (Raleigh, NC). Anti-goat IgG, anti-rat IgG, and anti-rabbit IgG antibodies were from Jackson ImmunoResearch (West Grove, PA). Enhanced chemiluminescence kit was obtained from Pierce (Rockford, IL), and Taq polymerase, from Roche Applied Science (Indianapolis, IN). Vectastain ABC kit, biotinylated anti-rabbit, anti-mouse IgG, or anti-rat IgG were from Vector Laboratories (Burlingame, CA). Trizol reagent, 2',7'-dichlorodihydrofluorescein diacetate (DCFDA), dihydroethidium (DHE), MitoSOX Red mitochondrial superoxide indicator, BLOCKIt Fluorescent Oligo, Lipofectamin, superscript II reverse transcriptase, 10–20% sodium dodecylsulfate (SDS), polyacrylamide gel, and 10–20% tricine gel were purchased from Invitrogen (Carlsbad, CA). Rac1 activation kit was purchased from Cell Biolab, Inc. (San Diego,

CA). Apocynin, paraquat, 3-(4,5-dimethylthiazal-2-yl)-2,5-diphenyl-tetrazolium bromide (MTT), protease inhibitor cocktail [AEBSEF, aprotinin, bestatin hydrochloride, E-64-[N-(trans-epoxysuccinyl)-L-leucine 4-guanidinobutylamide], leupeptin, pepstatin A], nitroblue tetrazolium (NBT) and dimethylsulfoxide (DMSO) were from Sigma-Aldrich (St. Louis, MO). CytoTox-96-NonRadioactive-Cytotoxicity-Assay for LDH activity was from Promega Bioscience (San Luis Obispo, CA). All other chemicals of reagent grade were from Sigma Chemicals or Merck (Rahway, NJ).

Animals and treatment paradigm

The experiments were carried out on mice, in accordance with the NIH *Guide for the Care and Use of Laboratory Animals*. All procedures were approved by the local Animal Care and Use Committee. Male C57BL/6 mice [Charles River (Wilmington, MA); 8 to 10 weeks] were maintained in a temperature/humidity-controlled environment under a 12-h light/dark cycle with free access to food and water. As depicted in Fig. 5A and B, each animal received three IP injections, separated by 2 days, of either vehicle (1% of DMSO in saline), PQ (10 mg/kg of body weight) or PQ combined with apocynin (200 mg/kg of body weight; Apo), according to previously published dosing (27, 40).

For the group co-treated with PQ and apocynin, an injection of apocynin was given before as well as on the day of the PQ injection. Five days after the last injection, animals were killed and were intracardially perfused for immunohistochemical analysis.

A time-course study (Fig. 5B) was also performed, in which the animals were intracardially perfused for immunohistochemical analysis 1 day after the first and second PQ injection and 5 days after the last injection. In this study, for the group co-treated with apocynin and PQ, apocynin was given before as well as on the day of the PQ injection, and the animals were killed 5 days after the last PQ injection (day 12).

In total, 38 mice (vehicle, PQ, and PQ+ Apo) were used in this study.

Cell-culture and treatment paradigm

The immortalized rat mesencephalic dopaminergic cell line (N27 cells) was grown in RPMI 1640 medium containing 10% FBS, 100 units of penicillin, and 50 μ g/ml streptomycin in a humidified atmosphere of 5% CO₂ at 37°C. For experiments, the cells were plated on polystyrene tissue-culture dishes at a density of 1 \times 10⁴ cells per well in 96-well culture plates, 0.5 \times 10⁵ cells per well in 24-well culture plates, 1.5 \times 10⁵ cells/well in six-well culture plates, or 5 \times 10⁵ cells per 100-mm dish. After 18 h, cells were treated with different concentrations of PQ or apocynin or both for the indicated duration. For siRNA transfection experiments, cells were plated at a density of 2 \times 10⁴ cells per well in 96-well culture plates and 5 \times 10⁵ cells per 60-mm dish.

Lactate dehydrogenase assay and MTT reduction assay

The extent of cell death was assessed by using the cytotoxic assay kit to evaluate the activity of lactate dehydrogenase (LDH) released into the culture medium. Aliquots (50 μ l) of cell-culture medium were incubated at room temperature in

the presence of 0.26 mM NADH, 2.87 mM sodium pyruvate, and 100 mM potassium phosphate buffer (pH 7.4) in a total volume of 200 μ l, for 15–30 min. The levels of NAD⁺ formation were measured at 490 nm by using a microplate spectrophotometer (SPECTRA MAX 340 pc; Molecular Devices, Sunnyvale, CA). To assess cell viability, levels of MTT reduction were measured. After 24-h PQ treatment, N27 cells were incubated with 0.5 mg/ml of MTT in medium overnight at 37°C. MTT is converted by viable cells to a water-insoluble precipitate that was dissolved in 10% SDS and colorimetrically quantified (O.D., 570 nm) by using a microplate spectrophotometer.

Determination of cellular ROS content

ROS levels were measured by using three different methods: DCFDA, DHE, and NBT assays. The DCFDA assay is based on the principle that the 2',7'-dichlorodihydrofluorescein diacetate can be oxidized by ROS and converted to the fluorescent 2',7'-dichlorofluorescein. In the DHE assay, blue fluorescent DHE can be dehydrogenated by superoxide (O₂⁻) to form a red fluorescent ethidium bromide. The NBT assay is based on the conversion of NBT to NBT diformazan (formazan dye) by superoxide radicals. PQ-treated cells were incubated with DCFDA (150 μ M), DHE (150 μ M), and NBT (0.3 mg/ml) in complete medium for 1, 4, and 6 h at 37°C, respectively. To measure the fluorescence produced in the DCFDA and DHE, the medium was removed, and PBS was added to each well. The emitted fluorescence was read in a microplate spectrophotometer plate reader at Ex/Em 502/535 or 525/620 nm for the DCFDA or DHE assays, respectively. To quantify NBT precipitation, cells were washed twice with 70% methanol and fixed for 5 min in 100% methanol. Wells were allowed to air dry, and the water-insoluble formazan was solubilized with 120 μ l 2 M KOH and 140 μ l DMSO. The optical density was read in a microplate spectrophotometer plate reader at 590 nm.

Western blot analysis

Cells were washed with ice-cold PBS and lysed on ice in RIPA buffer (50 mM Tris/HCl, pH 8.0, 150 mM NaCl, 2 mM sodium orthovanadate, 1% Nonidet-P40, 0.5% sodium deoxycholate, 0.1% SDS, containing 1% of a protease inhibitor mixture (AEBSE, pepstatinA, E-64, bestatin, leupeptin, and aprotinin) purchased from Sigma-Aldrich. The soluble fraction was obtained, and equal amounts of cell lysate protein were loaded in each lane of 10–20% SDS polyacrylamide gel. After electrophoresis and transfer onto a polyvinylidene difluoride membrane, specific protein bands were detected by using appropriate primary (rabbit anti-Nox1; mouse anti- β -actin) and secondary antibodies followed by enhanced chemiluminescence.

Rac1 activation assay

To assess Rac1 activation, we used the Rac Activation Assay Kit, from Cell Biolabs, Inc. (SKU STA-401; San Diego, CA). We followed the protocol provided by the manufacturer. In brief, N27 cells were treated with 500 and 800 μ M PQ for 8 h, followed by total cellular proteins extraction. Of the total cellular proteins, 1 mg was incubated with 10 mg of agarose

beads containing the p21-binding domain (PBD) of the p21-activated protein kinase 1 (PAK1), an effector of activated Rac, for 1 h at 4°C. The beads were collected by centrifugation and washed two times in the lysis buffer. The beads were resuspended in sample buffer and boiled for 5 min. Proteins were resolved by SDS-PAGE with a 10–20% Tricine gel, transferred electrophoretically, and visualized by using anti-rat Rac1 antibody followed by enhanced chemiluminescence. For the positive control, the nonhydrolyzable GTP analogue GTP γ S was used according to the manufacturer's protocol.

Preparation and transfection of siRNA

Sense and antisense oligonucleotides corresponding to the rat Nox1 cDNA sequences 5'-CCTTTGCTTCCTTCTTGAAATCTAT-3', which does not contain any sequences homologous to other Nox enzymes, were used. The double-stranded siRNAs were synthesized chemically and modified into stealth siRNA to enhance the stability *in vitro*. As negative control, stealth siRNA with a GC content similar to that of Nox1 stealth siRNA was used. When N27 cell cultures reached ~80% confluence in 96-well culture plates or 60-mm dishes, transfection by adding Lipofectamin 2000 and siRNAs (final concentration, 33 nM) was performed. After 6 hr of incubation, the culture medium was changed, and cells were maintained for additional 36 h before PQ treatment. To evaluate transfection efficiency, N27 cells were also transiently transfected with siRNA tagged with fluorescence (BLOCKiT Fluorescent Oligo) for 36 h, and transfected cells were identified by green fluorescence in cells. Double-stranded siRNAs, the negative control stealth siRNA, as well as the BLOCKiT Fluorescent Oligo, were purchased from Invitrogen.

Immunohistochemistry

After perfusion with saline and 4% paraformaldehyde in phosphate-buffered saline (PBS), brains were removed, and forebrain and midbrain blocks were immersion-fixed in 4% paraformaldehyde and cryoprotected in sucrose. Serial coronal sections (40 μ m) were cut on a cryostat, collected in cryopreservative, and stored at -20°C. For immunolabeling studies, sections were incubated with blocking solution (5% horse serum and 0.3% Triton X-100 in PBS, pH 7.5) and then with primary antibodies at room temperature overnight. Finally, sections were incubated with secondary antibodies in blocking solution at room temperature for 1 h. The primary antibodies used were rabbit anti-TH antibody (1:10,000; Protos Biotech, New York, NY), rabbit anti-Nox1 (1:500; Santa Cruz Biotechnology, rat anti-CD11b (1:50; Serotec, Raleigh, NC). The secondary antibodies were biotinylated anti-rabbit IgG or anti-rat IgG (1:200; Vector Laboratory and following the staining procedure outlined by the manufacturer of the Vectastain ABC kit in combination with 3,3'-diaminobenzidine reagents. The same TH-stained sections were then counterstained with Nissl (0.25% cresyl violet) for 5 min, washed in distilled water, air dried, cleaned in xylene, and then mounted with the appropriate mounting medium. The numbers of TH-immunoreactive cells and Nissl-positive cells in the SN pars compacta (SNpc) were counted by using an optical fractionator. Analysis was performed by using a system consisting of a Nikon Eclipse E600 microscope (Morrell Instruments Co. Inc., Melville, NY) equipped with a computer-controlled

LEP BioPoint motorized stage (Ludl Electronic Products, Hawthorne, NY), a DEI-750 video camera (Meyer Instruments, Houston, TX), a Dell Dimension 4300 computer (Dell, Round Rock, TX), and the Stereo Investigator (v. 4.35) software program (Microbrightfield, Burlington, VT). Tissue sections were examined by using a Nikon Plan Apo 100x objective lens with a 1.4 numeric aperture. The size of the x-y sampling grid was 140 μm . The counting-frame thickness was 14 μm , and the counting-frame area was 4,900 μm^2 .

Total RNA extraction and RT-PCR analysis

Total RNA was extracted from N27 cells by using Trizol reagent. Reverse transcription (RT) was performed for 40 min at 42°C with 1 μg of total RNA by using 1 unit/ μl of superscript II reverse transcriptase. Oligo (dT) and random primers were used as primers. The samples were then heated at 94°C for 5 min to terminate the reaction. The cDNA obtained from 1 μg total RNA was used as a template for PCR amplification. Oligonucleotide primers were designed based on Genebank entries for rat Nox1 (sense, 5'-TGACAGTGATGTATGCAGCAT-3'; antisense, 5'-CAGCTTGTTGTGTGCACGCTG-3'), rat Nox2 (sense, 5'-ACTCGAAAACCTTCTTGGGTCAG-3'; antisense, 5'-TCCTGTGATGCCAGCCAACCGAG-3'), rat Nox3 (sense, 5'-GCTGGATTTTGAACGAGAGTGTG-3'; antisense, 5'-GCCAGAGAGATCACCAGGCCAGT-3'), rat Nox4 (sense, 5'-GCCGGCGGTATGGCGCTGTC-3'; antisense, 5'-CCACCATGCAGACACCTGTCAGG-3'), and rat GAPDH (sense, 5'-ATCACCATCTTCCAGGAGCG-3'; antisense, 5'-GATGGCATGGACTGTGGTCA-3'). PCR mixes contained 10 μl of 2x PCR buffer, 1.25 mM of each dNTP, 10 pmol each of forward and reverse primers, and 2.5 units of Taq polymerase to a final volume of 20 μl . For Nox1, amplification was performed in 32 cycles for 40 s at 95°C, 30 s at 62°C, and 2 min at 72°C, and for Nox2, Nox3, and Nox4, it was performed in 27 cycles for 40 s at 95°C, 30 s at 58°C, and 2 min at 72°C. After the last cycle, all

samples were incubated for an additional 7 min at 72°C. PCR fragments were analyzed on 1% agarose gel containing ethidium bromide, and their amounts were normalized against amplified GAPDH. Each primer set specifically recognized only the gene of interest, as indicated by amplification of a single band of the expected size.

Data analysis and statistics

Data are expressed as percentages of values obtained in control conditions and are presented as mean \pm SEM of at least three experiments, in independent cell cultures. Statistical analysis was performed by using a one-way ANOVA followed by the Dunnett test or the Bonferroni Multiple Comparison Test. Values of $p < 0.05$ were considered significant.

Results

Paraquat-mediated ROS generation and subsequent dopaminergic cell death

It is known that PQ directly generates superoxide through redox cycling. To evaluate the effect of this compound on ROS generation, we treated N27 cells with various concentrations of PQ (100, 500, 800, or 1,000 μM) for 24 h and measured ROS levels with NBT, DCFDA, and DHE assays. The results obtained by using the NBT assay showed that cultures treated with 100 μM PQ showed $34 \pm 2.3\%$ higher levels of ROS than did control cultures. The levels of ROS were further increased in cultures treated with higher concentrations of PQ: $70 \pm 4.5\%$ increase in cultures treated with 500 μM , $65 \pm 4.6\%$ increases with 800 μM , and $111 \pm 7.2\%$ increase with 1,000 μM of PQ (Fig. 1A). When ROS was measured with DCFDA (Fig. 1B), similar changes were observed ($46 \pm 13\%$, $53 \pm 7\%$, and $82 \pm 11\%$ increase in 500, 800, and 1,000 μM PQ, respectively),

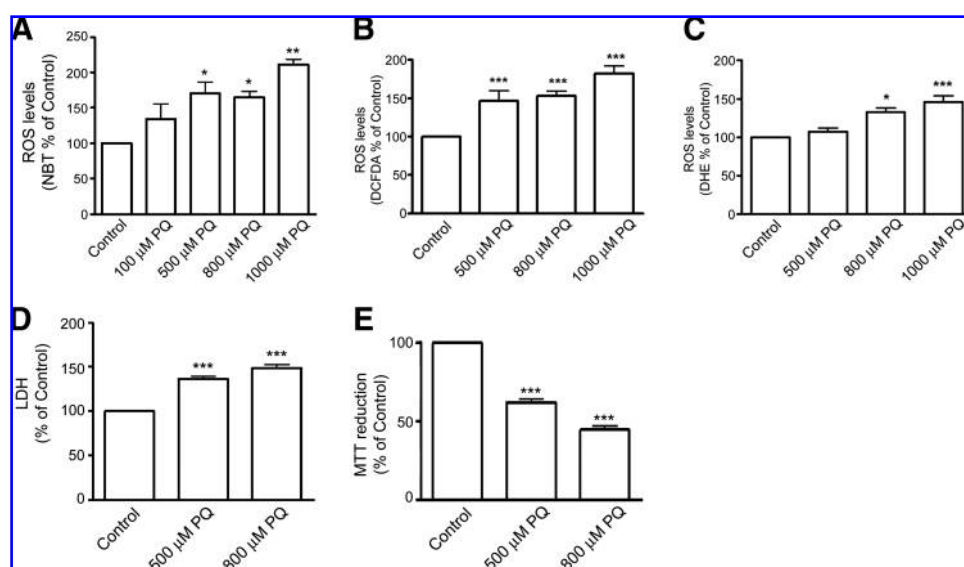


FIG. 1. PQ-mediated ROS generation and N27 DA cell death in a concentration-dependent manner. (A) ROS levels in N27 cells treated with PQ (100, 500, 800, or 1,000 μM) for 24 h was measured by using the NBT assay (A). ROS levels in N27 cells treated with PQ (500, 800 or 1,000 μM) for 24 h measured by using DCFDA (B) or DHE (C). (D, E) Cell-death extent was accessed by measuring the levels of LDH activity (D) and MTT reduction (E) in N27 cultures incubated with 500 or 800 μM PQ for 24 h. The results are expressed as percentage of control. Data are shown as the mean \pm SEM of three independent experiments performed in triplicate. Statistical

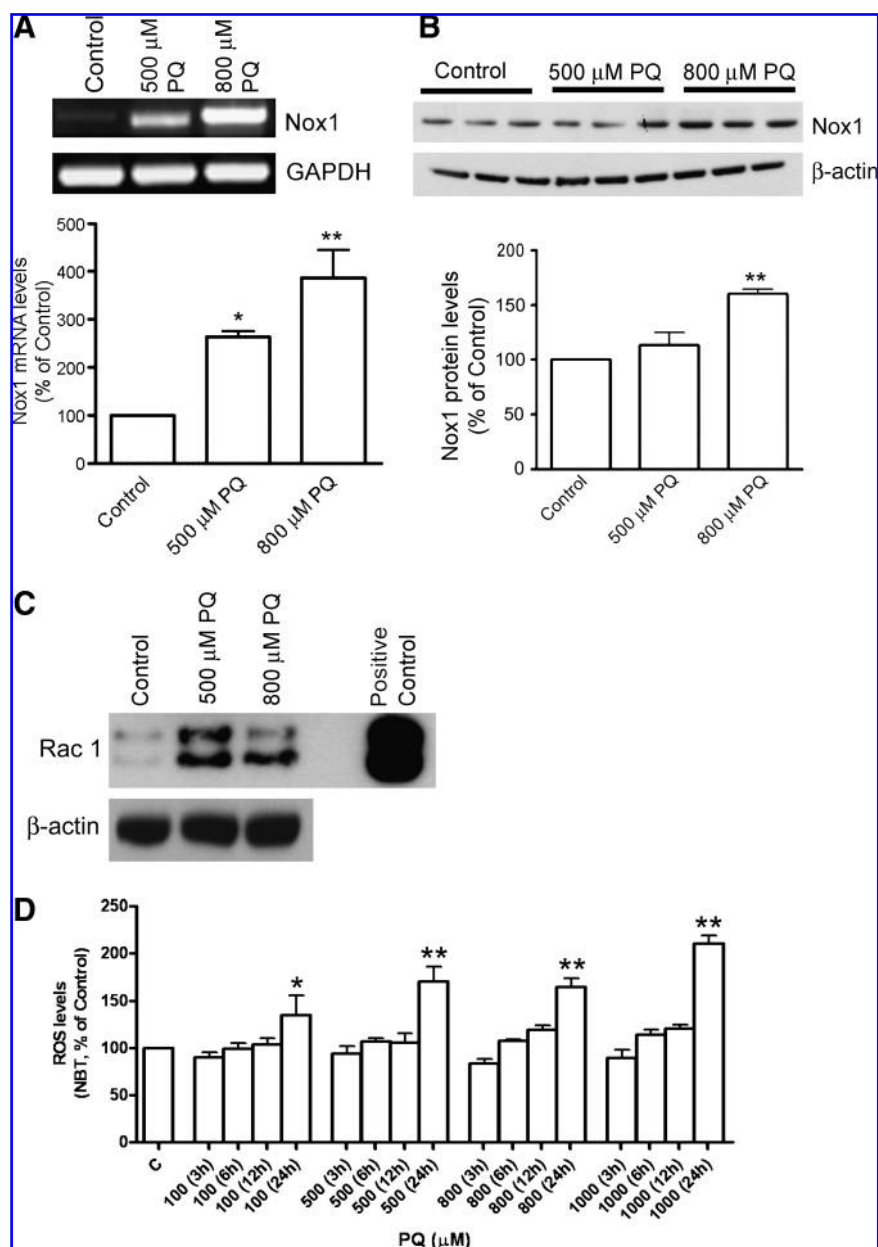
analysis was performed by using one-way ANOVA followed by Bonferroni's Multiple Comparison Test. * $p < 0.05$, ** $p < 0.01$, and *** $p < 0.001$ versus control cultures.

whereas the DHE assay (Fig. 1C) showed that significant superoxide increase was elicited only by 800 and 1,000 μM PQ ($33 \pm 5\%$ and $45 \pm 7\%$ increase compared with control, respectively). To evaluate the effect of PQ on DA cell death, we investigated the levels of LDH and reduced MTT in N27 cells after treatment with 500 or 800 μM PQ for 24 h. As depicted in Fig. 1D, LDH levels were increased by $36 \pm 4\%$ in cultures treated with 500 μM PQ and increased by $49 \pm 6\%$ in cultures treated with 800 μM , in comparison with control cultures. Similar results were obtained in MTT assays measuring viable cells that are capable of reducing MTT. Cultures treated with 500 μM PQ showed $62 \pm 1.4\%$ of the cell viability, and only $44 \pm 2.8\%$ cells survived in cultures treated with 800 μM PQ (Fig. 1E). As demonstrated in a previous study (45), these results showed that PQ elevates ROS levels in N27 cells and finally leads to DA cell death in a concentration-dependent manner.

Paraquat-mediated increase in Nox1 expression and the activation of Rac1 in dopaminergic cells

The Nox system catalyzes the reduction of molecular oxygen and oxidation of NADPH to generate superoxide radicals and, in this way, contributes to PQ redox cycling and eventually to its toxicity. We investigated whether DA cells have the NADPH oxidase enzyme complex and whether it is affected by PQ. We evaluated mRNA levels of Nox1 in N27 DA cells treated with 500 or 800 μM PQ for 6 h, as well as in nontreated controls. Interestingly, we observed that mRNA levels of Nox1, the first Nox2 homologue identified in non-phagocytic cells, were significantly increased by PQ in a concentration-dependent manner (Fig. 2A, upper panel). N27 cells had all components of Nox1 activation, including Rac1, Noxa1, and Noxo1 (data not shown). Cultures treated with 500 μM PQ showed a $163 \pm 13\%$ increase in Nox1 mRNA

FIG. 2. Increased Nox1 expression levels and the activation of Rac1 in N27 DA cells treated with PQ. (A) Nox1 mRNA level was detected with RT-PCR (upper panel), in N27 cells treated with 500 or 800 μM PQ for 6 h, and quantified with Quantity One software and then normalized to GAPDH, an internal control (lower panel). (B) Nox1 protein levels were obtained with immunoblot analysis in N27 cells treated with 500 or 800 μM PQ for 16 h (upper panel) and quantified by using Quantity One software (lower panel). β -Actin levels were measured as an internal control. (C) The activated fraction of Rac1 was determined in N27 cells treated with 500 or 800 μM PQ for 8 h with the active GTPase pull-down assay. β -Actin levels were measured as an internal control. (D) The time-course measurement of PQ-mediated ROS generation in N27 cells. ROS levels in N27 cells treated with PQ (100, 500, 800 or 1,000 μM) for 3, 6, 12 and 24 h were measured by using the NBT assay. The results are expressed as percentage of control. Data are shown as the mean \pm SEM of three independent experiments. Statistical analysis was performed by using one-way ANOVA followed by Bonferroni's Multiple Comparison Test for Nox1 mRNA levels and by Dunnett's Multiple Comparison Test for Nox1 protein levels. * $p < 0.05$; ** $p < 0.01$ versus control cultures.



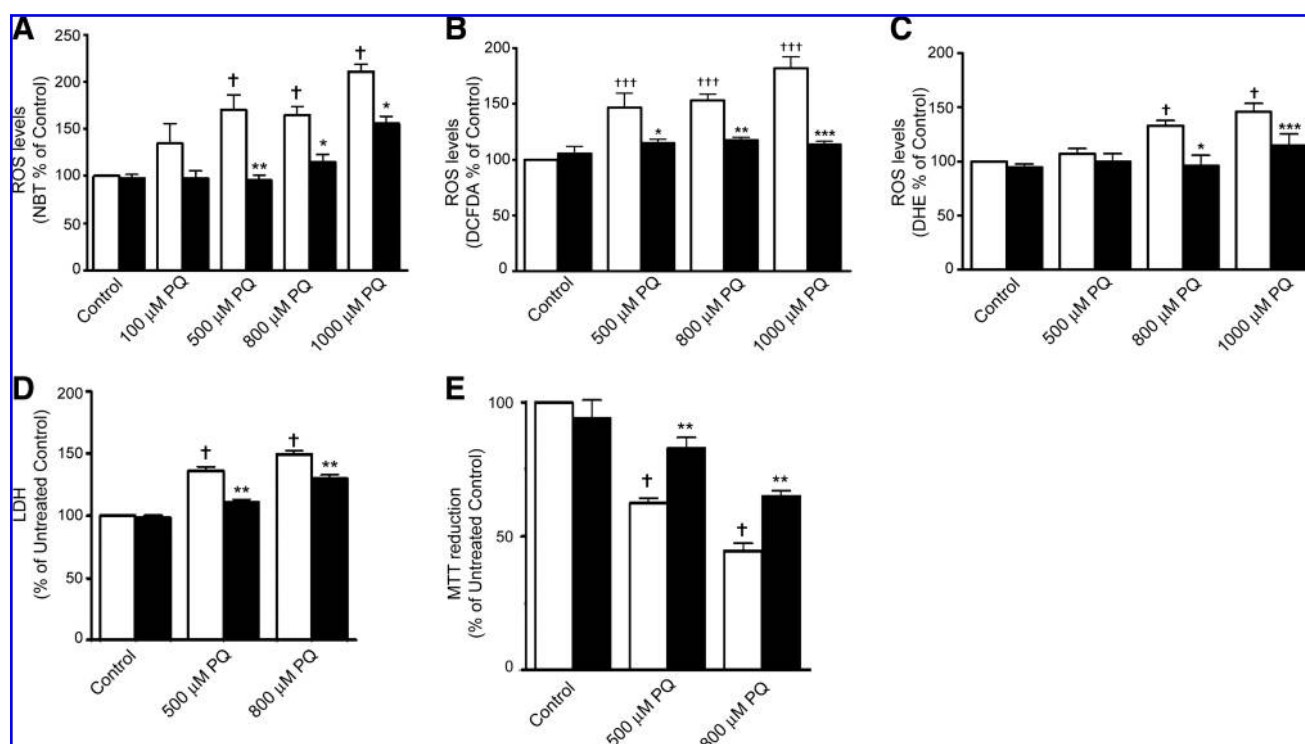


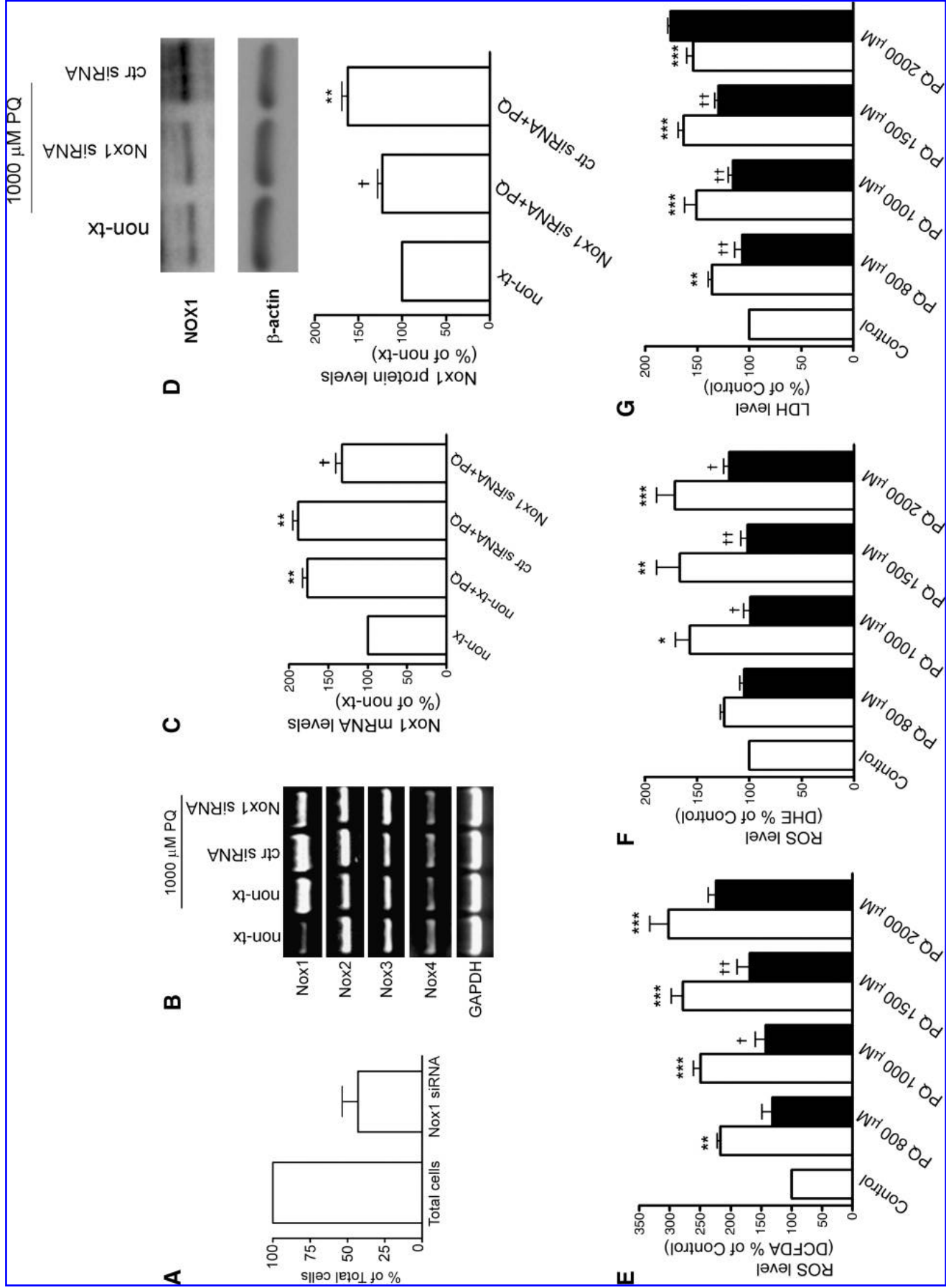
FIG. 3. Decreases in PQ-mediated ROS levels and N27 DA cell death by a putative NADPH oxidase inhibitor, apocynin (A–C). Apocynin significantly reduced PQ-mediated ROS generation by N27 cells. ROS levels were measured by using the NBT (A) assay, DCFDA (B), or DHE (C) in N27 cells pretreated with 5 μ M apocynin for 1 h and then treated with various concentrations of PQ (100, 500, 800, or 1,000 μ M) for 24 h. (D, E) Apocynin significantly reduced PQ-mediated N27 cell death. Cell-death levels were assessed by measuring the LDH activity (D) and MTT reduction (E) in N27 cultures pretreated with 5 μ M apocynin for 1 h and then treated with 500 or 800 μ M PQ for 24 h. The results are expressed as percentage of their controls. Data are shown as the mean \pm SEM of three independent experiments performed in triplicate. Statistical analysis was performed by using one-way ANOVA followed by Bonferroni's Multiple Comparison Test. $^*p < 0.05$; $^{**}p < 0.01$; and $^{***}p < 0.001$ versus cultures pretreated with apocynin and then with PQ. $^{\dagger}p < 0.05$, and $^{\dagger\dagger\dagger}p < 0.001$ versus control cultures. Solid bars, cells pretreated with apocynin; open bars, cells without apocynin treatment.

levels compared with control cultures, and higher PQ doses (800 μ M) increased Nox1 mRNA levels to $287 \pm 62\%$, compared with controls (Fig. 2A, lower panel). Next, we evaluated Nox1 protein levels in N27 cells incubated for 16–18 h with or without PQ. Only cultures treated with 800 μ M PQ showed a significant increase in Nox1 protein levels ($60 \pm 6\%$; Fig. 2B). Although 500 μ M PQ was able to increase Nox1 mRNA (Fig.

2A), this concentration was not sufficient to increase significantly Nox1 protein levels ($13 \pm 10\%$) compared with controls (Fig. 2B). The functional activation of Nox1 is dependent on the activation of a small GTPase, Rac1 (9).

Based on these facts, we tested whether PQ induces the activation of Rac1. Activated Rac1 was limited in nontreated N27 cells. After treatment of cells with PQ for 8 h, a significant

FIG. 4. Decreases in PQ-mediated ROS levels and N27 DA cell death by Nox1 knockdown. (A) Transfection efficiency of Nox1 siRNA into N27 cells. (A) N27 cells were transiently transfected with siRNA tagged with fluorescence for 36 h. Transfected cells were identified with green fluorescence in cells, and total cells were identified with DAPI staining. Transfection efficiency was calculated as percentage of green fluorescence-positive cells to total cells stained for DAPI. Nox1 siRNA-mediated knockdown efficiency was also verified by using RT-PCR for Nox1, Nox2, Nox3, and Nox4 (B and C) and immunoblot analysis for Nox1 (D). N27 cells were transiently transfected with rat Nox1 siRNA, control siRNA (ctr siRNA), or exposed only to lipofectamine (non-tx) for 36 h and then treated with 1,000 μ M PQ. GAPDH and β -actin were used as internal controls for RT-PCR and immunoblot, respectively. Nox1 mRNA (C) and protein (D) levels were quantified by using Quantity One software from Bio-rad. For C and D, $^{**}p < 0.01$ versus non-tx and $^{\dagger}p < 0.05$ versus ctr siRNA+PQ or versus non-tx+PQ or both. (E–G) Nox1 knockdown significantly reduced PQ-mediated ROS levels as well as N27 cell death. ROS levels were measured by using DCFDA (E) or DHE (F), and cell death was evaluated by using the LDH assay (G) in N27 cells transiently transfected with rat Nox1 siRNA and then treated with various concentrations of PQ (800, 1,000, 1,500, or 2,000 μ M) for 24 h. $^*p < 0.05$; $^{**}p < 0.01$; and $^{***}p < 0.001$ versus control. $^{\dagger}p < 0.01$ and $^{\dagger\dagger}p < 0.05$ versus ctr siRNA+PQ. Solid bars, cells transfected with Nox1 siRNA; open bars, cells transfected with ctr siRNA. Data are shown as the mean \pm SEM of three independent experiments performed in triplicate. Statistical analysis was performed by using one-way ANOVA followed by Bonferroni's Multiple Comparison Test.



increase in activated Rac1 was observed (Fig. 2C). To investigate a sequential relation between Nox1 activation and PQ-mediated ROS generation, the time-course of ROS generation was investigated in N27 cells treated with various doses of PQ (Fig. 2D). ROS levels were significantly increased only after 24-h PQ treatment, suggesting that Nox1 activation can be responsible for PQ-mediated ROS generation.

Attenuation of PQ-mediated ROS generation and dopaminergic cell death by inhibition of NADPH oxidase

Apocynin is a putative inhibitor of NADPH oxidase activity. To clarify the specific role of NADPH oxidase in PQ-mediated ROS generation and DA cell death, we tested the effect of apocynin in N27 cells treated with PQ. N27 cells were pretreated with 5 μ M apocynin for 1 h, and then different concentrations of PQ (100, 500, 800, or 1,000 μ M) were added. ROS levels were measured at 24 h after PQ by using an NBT assay. Apocynin significantly reduced PQ-mediated ROS generation in N27 cells (Fig. 3A). ROS levels in cells treated with 500 and 800 μ M PQ were significantly reduced compared with control levels. Apocynin was also able to significantly reduce ROS levels induced by relatively high dose of PQ (1,000 μ M) by $55 \pm 5\%$. ROS levels measured by using DCFDA (Fig. 3B) or DHE (Fig. 3C) showed similar changes observed by using the NBT assay. The effect of apocynin on PQ-mediated DA cell death was also tested. Cultures were pretreated with 5 μ M apocynin for 1 h and then treated with 500 or 800 μ M PQ. The viability of cells was determined with LDH and MTT assays (Fig. 3D and E). Apocynin significantly reduced LDH levels by $26 \pm 7\%$ and $19 \pm 5\%$ in cultures treated with 500 μ M and 800 μ M PQ, respectively, compared with cultures treated with PQ alone (Fig. 3D). The levels of reduced MTT were statistically higher in cultures pretreated with apocynin than in those treated with PQ only. As shown in Fig. 3E, apocynin increased the levels of reduced MTT by $21 \pm 5\%$ and $20 \pm 4\%$ in cultures treated with 500 μ M and 800 μ M PQ, respectively.

Even though apocynin has been widely used in several studies as an inhibitor of the NADPH oxidase system (8, 27), recent studies also suggest that this compound may work as a direct antioxidant (3, 29). Thus, to investigate the specific role of Nox1 in PQ-mediated oxidative stress and DA cell death, siRNA-mediated Nox1 knockdown was used as an alternative tool. Transfection efficiency of siRNA nucleotide sequence into N27 cells was evaluated by using fluorescent-tagged siRNA: $42 \pm 10\%$ of total cells were efficiently transfected after 36-h incubation (Fig. 4A). Accordingly, PQ-induced mRNA (Fig. 4B and C) levels of Nox1 were significantly reduced compared with nontransfected ($43 \pm 6\%$ less) and control siRNA transfected cells ($55 \pm 5\%$ less). PQ-induced protein (Fig. 4D) levels of Nox1 were also significantly reduced ($38 \pm 7\%$) compared with cells transfected with control siRNA. To ensure selective knockdown of Nox1 by siRNA, mRNA levels of other Nox isoforms, including Nox2, Nox3 and Nox4, were determined as well. No changes were found between nontransfected cells and cells transfected with control siRNA or Nox1 siRNA (Fig. 4C). Finally, PQ-mediated ROS generation was also significantly decreased in cells transfected with Nox1 siRNA (Fig. 4E and F). PQ-mediated DA cell death was also

significantly reduced by Nox1 knockdown (Fig. 4G). Although Nox1 knockdown significantly inhibited ROS generation by 2,000 μ M PQ, cell death was not prevented at this concentration, implying that PQ induces cell death through pathways distinct from ROS-mediated pathways.

These results highlight that the NADPH oxidase system, and specially Nox1, plays a crucial role in PQ-mediated ROS generation as well as subsequent DA cell death.

Increased expression of Nox1 in the SN of mice treated with PQ intraperitoneally

Previous *in vivo* studies tested various PQ-injection paradigms in mice and demonstrated a selective DA neuronal death (41, 42, 49). An approximate 25% DA cell loss was reported in mice treated with 10 mg/kg IP of PQ every week for 3 weeks (42). To ensure significant levels of PQ toxicity on DA neurons, we tested a different injection paradigm, depicted in Fig. 5A. Each animal received three IP injections, separated by 2 days of either vehicle (1% of DMSO in saline), PQ (10 mg/kg), or PQ combined with apocynin (200 mg/kg). Apocynin was also administered IP 1 day before and on the same day as PQ injection. A similar dose of apocynin (300 mg/kg supplied by drinking water) showed substantially increased animal life span in ALS mice with mutant SOD1 (G93A) (27). All groups were killed 5 days after the last injection. This injection protocol resulted in about a $35 \pm 8\%$ reduction of TH-positive neurons in the SNpc compared with that in animals treated with vehicle (Fig. 7A and B). In comparison with animals treated with vehicle, those treated with PQ showed a slight decrease in body weight and overall activity. After the last injection, animals gradually recovered over the last 5 days of the experiment (data not shown). Mice treated with PQ and apocynin were more active and lost less weight, when compared with mice treated with PQ only.

We also tested another injection paradigm in which mice were treated with three injections of a higher dose of PQ (15 mg/kg) separated by 1-day intervals. At this high-dose and high-frequency paradigm, all animals died after the second injection, suggesting its systemic toxicity. These results indicate that PQ has a narrow window in which selective DA neuronal death is observed without producing overt systemic toxicity.

To verify our *in vitro* finding showing increased Nox1 levels in DA cells insulted by PQ, Nox1 levels in the SN of mice treated with vehicle or PQ were investigated with Western blot and RT-PCR analysis. As shown in Fig. 6, animals treated with PQ showed statistically higher levels of the Nox1 protein than did the animals treated with vehicle (Fig. 6A and B). Nox1 mRNA was also increased (Fig. 6C). Upregulation of Nox1 in the SN of mice injected with PQ was confirmed by immunostaining (Fig. 6D).

Apocynin reduced DA neuronal death induced by PQ in mice

Each group of animals was injected as described earlier, and the numbers of TH-positive cells and Nissl-positive cells in the SNpc were stereologically counted. As shown in Fig. 7, PQ administration led to a $35 \pm 13.7\%$ reduction of TH-

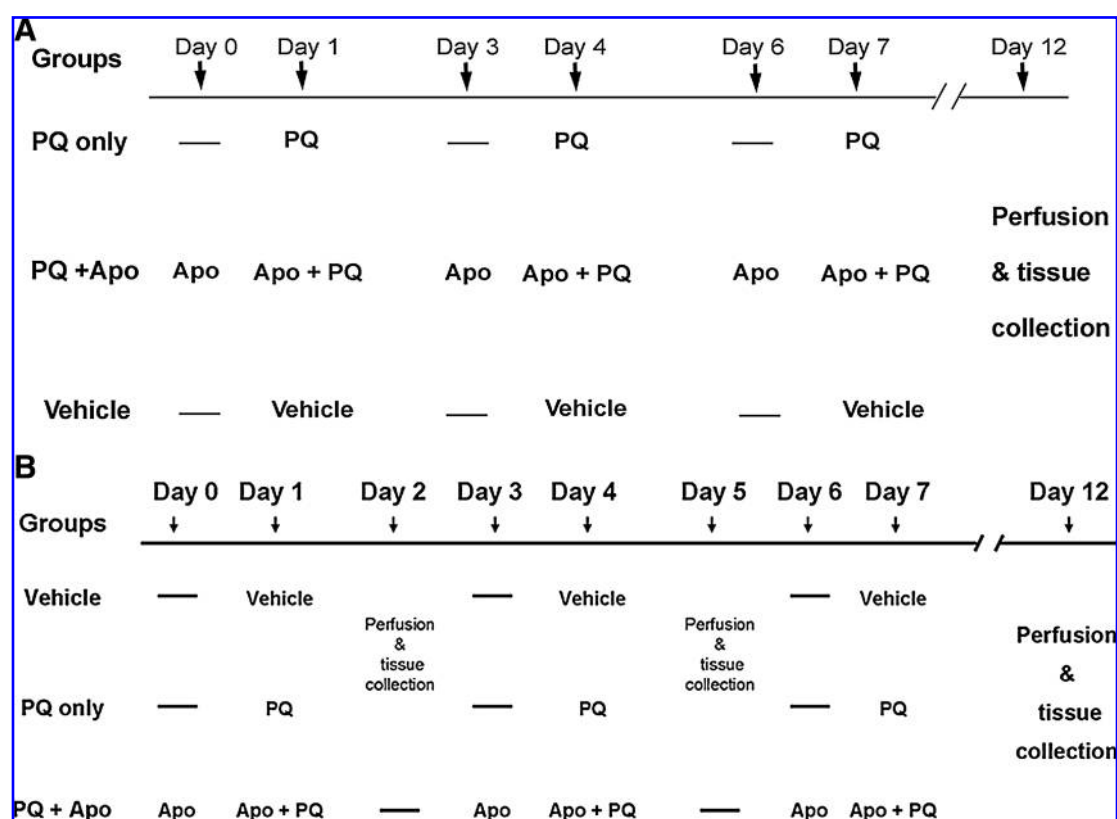


FIG. 5. PQ-injection paradigm diagram. Mice were divided into three groups. Animals were given a total of three injections of either vehicle (1% DMSO in saline), PQ (10 mg/kg of body weight), or PQ combined with apocynin (200 mg/kg of body weight) every third day. Apocynin was also administered IP 1 day before and on the same day as PQ injection. All groups were killed 5 days after the last injection (**A**); to investigate PQ-mediated microglia changes, animals were killed the day after the first and second injections and 5 days after the last injection (**B**).

positive neurons in the SNpc compared with those in animals treated with vehicle (Fig. 7A and B). Apocynin administration significantly reduced PQ-elicited TH⁺ neuronal loss to $15.7 \pm 3.8\%$. Total neuronal number in the SN by Nissl staining also indicated decreased neuronal staining in the SNpc of PQ-treated mice ($25.3 \pm 6.5\%$) and that the combined treatment with apocynin elevated neuronal numbers in the SNpc to $91.3 \pm 7.8\%$ (Fig. 7A and C).

Our results support the idea that the NADPH oxidase system plays an important role in PQ-mediated DA neurotoxicity.

Paraquat-mediated microglial changes

It was previously reported that PQ exerts its toxic effects on DA neurons through microglial activation (43, 47, 59). Thus, the time-course morphologic changes in microglia in the SN after PQ treatment (Fig. 5B) were investigated with CD11b immunostaining at days 2, 5, and 12 after the first PQ injection (Fig. 8). Whereas vehicle-treated animals showed the typical ramified morphology of resting microglia, PQ treatment resulted in dramatic morphologic changes in microglia, characterized by enlarged cell bodies and loss of processes. These morphologic changes occurred at an early time point (2 days). The total number of CD11b-positive microglia was gradually decreased thereafter and was not detected at day 12 (Fig. 8A

and B). Similar changes were observed in animals treated with apocynin in combination with PQ (Fig. 8A).

Discussion

In this study, we showed that the Nox complex mediates oxidative stress and cell death caused by paraquat in N27 dopaminergic cells. We identified for the first time that the Nox1 isoform is constitutively expressed in DA cells, and its level is elevated by paraquat administration both *in vivo* and *in vitro*. NADPH oxidase-derived superoxide generation and bacterial killing were first discovered in polymorphonuclear neutrophils (Nox2 or gp91^{phox}) (50). Nox1 is the first homologue of Nox2 that was identified in nonphagocytic cell types, including colon, prostate, uterus, and vascular smooth muscles (55). Since its discovery, a family of homologues has been identified in a variety of cells and tissues. Previous studies have shown that Nox isoforms are expressed in mouse and rat brain tissues (36, 52). In the CNS, although NADPH oxidase-mediated ROS are required for normal cellular functions, such as long-term potentiation (57) and cardiovascular homeostasis (48), excess ROS generation may contribute to pathologic conditions. Oxidative stress elicited by Nox2 in microglia, an immune component in the brain, has been widely studied in a number of brain diseases (16, 37). Recent studies, however, indicate that Nox2 expression is not limited to microglia (1),

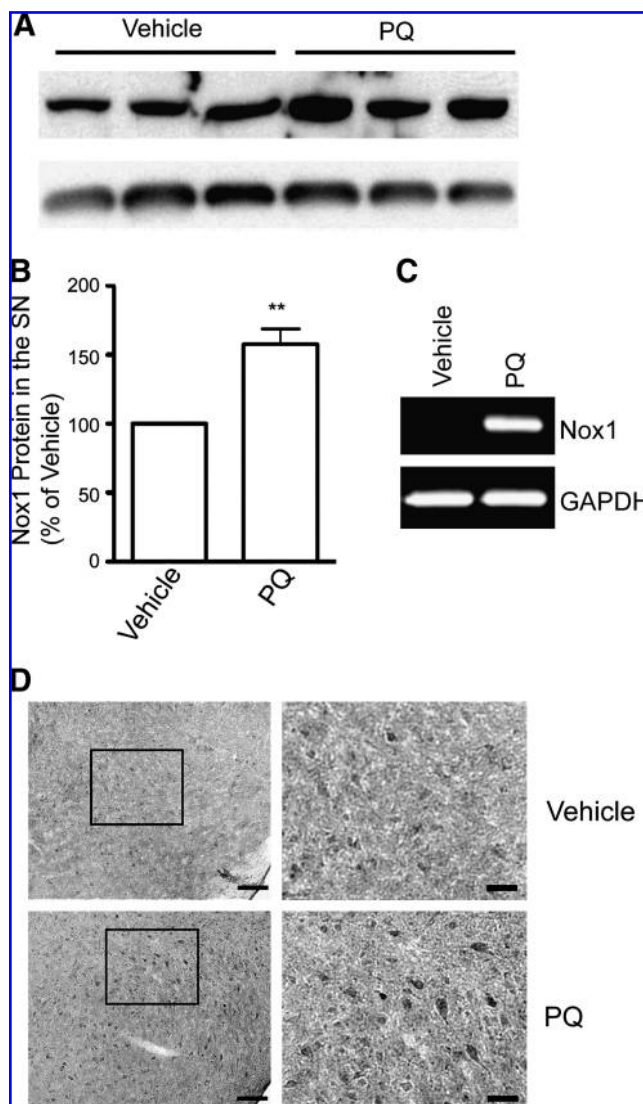


FIG. 6. Significant increase in Nox1 protein levels in the SNpc of mice injected with PQ. (A) Nox1 protein levels were determined in total lysates of the SN tissues of mice injected with PQ or vehicle with immunoblot analysis and RT-PCR. β -Actin and GAPDH were detected as internal control. PQ significantly increased Nox1 protein (A, B) and mRNA levels (C). Nox1 protein levels were quantified by using Quantity One software and normalized against β -actin (B). The results are expressed as percentage of vehicle. Data are shown as the mean \pm SEM of three independent experiments ($n=3$). Statistical analysis was performed by using one-way ANOVA followed by Bonferroni's Multiple Comparison Test. $**p < 0.01$ versus mice treated with vehicle. (D) Representative photomicrograph of Nox1-immunoreactivity in the SNpc sections of mice injected with vehicle or PQ. Nox1 immunoreactivity in the SN was increased in PQ-injected animals compared with vehicle. Right panels of D show higher magnification of respective boxed areas in the left panels. Scale bar in the left panels = 100 μ m, and in the right panels, 50 μ m.

and other homologues, including Nox1, are involved in various pathologic conditions in neuronal cells. Glutamate toxicity in SH-SY5Y neuroblastoma cells is largely attenuated by inhibition of NADPH oxidase activation (44). ROS generation

and apoptosis of N27 DA cells treated with MPP⁺, an active metabolite of MPTP, was decreased by Nox inhibition in one study (4). In the study, however, the authors did not clearly demonstrate which isoforms are responsible for MPP⁺-mediated ROS generation.

Our results demonstrated that MPP⁺ increased Nox1 expression in N27 cells (data not shown), suggesting that Nox1-mediated oxidative stress might be a common feature observed in DA cells under toxic stimuli. An increasing number of studies revealed that Nox1 plays a variety of roles in various neuronal cells. Nox1-derived superoxide negatively affects NGF-mediated neurite outgrowth in PC12 cells (31). In dorsal root ganglion cells, ROS generated from Nox1 are involved in enhancing sensitivity to painful stimuli (32). Increased transcription activity of Nox1 was observed in N27 cells treated with PQ in a concentration-dependent manner. Interestingly, several studies suggested that a crosstalk between mitochondria and Nox1 induction may exist. In 293 cells, mitochondrial ROS generated under serum deprivation induces Nox1 expression and superoxide generation. Although mitochondria contribute to the early accumulation of ROS (0–4 h), the maintenance of ROS level requires PI3K/Rac1/Nox1 activation. This later phase of ROS generation and sustained accumulation by Nox1 is responsible for cell death (38). These results suggest that although the initial stage of ROS generation is responsible for triggering the Nox system, sustained superoxide production from this system is an essential component of ROS-induced cell death.

Studies of osteosarcoma cells that lack a mitochondrial genome (ρ^0) revealed that the inactivation of mitochondrial genes leads to the downregulation of Nox1 and that transfer of wild-type mitochondrial genes can restore Nox1 expression (19). Our results also confirmed that mitochondrial respiratory-chain inhibitors (including rotenone, pyridaben, antimycin A, and FCCP) elevated both mRNA and protein levels of Nox1 (data not shown). Several molecular pathways have been suggested to understand PQ-mediated oxidative stress. Despite its structural similarity with MPP⁺, each compound has distinct molecular mechanisms for exerting oxidative stress and cellular damage (49). PQ is capable of directly generating superoxide through redox cycling (11), by which it may accept electrons from NAD(P)H and subsequently reduce molecular oxygen to superoxide. Mitochondrial respiratory complexes I and III both serve as targets for PQ-mediated ROS generation, with complex III showing a higher sensitivity (12, 49). Taken together, PQ-induced increase in Nox1 transcriptional activity might result from initial ROS produced by the PQ redox cycle or mitochondria. PQ may also activate signaling pathways such as PKC delta or MAPK, which were reported as Nox1 transcriptional activators (13, 23).

Nox1-mediated superoxide generation requires other components, including Rac1 activation, Nox1 and Noxa1, and homologues of p47^{phox} and p67^{phox}, respectively (15). Nox1 and Noxa1 mRNAs were constitutively expressed at high levels (data not shown). Rac1, a small Rho family GTPase, is an important subunit for the activation of Nox1-derived superoxide generation (15, 58). To activate the NADPH oxidase system, activated Rac1 forms a Nox1 enzyme complex in conjunction with Nox1 and Noxa1. In the present study, PQ-mediated Rac1 activation was observed.

FIG. 7. Apocynin reduced SNDA neuronal death induced by PQ in a mouse PD model.

(A) Representative photomicrographs of TH-immunostaining and Nissl counterstaining in the SNpc of mice treated with vehicle, PQ alone, or PQ and apocynin. Scale bar in the middle panel, 200 μ m. (B) Changes in the number of TH-immunoreactive neurons in the SNpc after PQ treatment in the presence or absence of apocynin. The number of TH-positive neurons was statistically decreased in animals treated with PQ. This decrease was significantly attenuated in mice co-treated with PQ and apocynin. (C) Changes in the numbers of total neurons (Nissl staining) in the SNpc after PQ treatment in the presence or absence of apocynin. The number of total neurons was statistically decreased in animals treated with PQ. This decrease was significantly attenuated in mice co-treated with PQ and apocynin. The results are expressed as percentage of vehicle. Data are shown as the mean \pm SEM (vehicle group, $n=6$; PQ group, $n=7$; Apo+PQ group, $n=5$). Statistical analysis was performed by using one-way ANOVA followed by Bonferroni's Multiple Comparison Test. * $p < 0.05$ and *** $p < 0.001$ versus mice treated with vehicle; $^{\dagger}p < 0.05$ and $^{\dagger\dagger}p < 0.01$ versus mice treated with PQ.

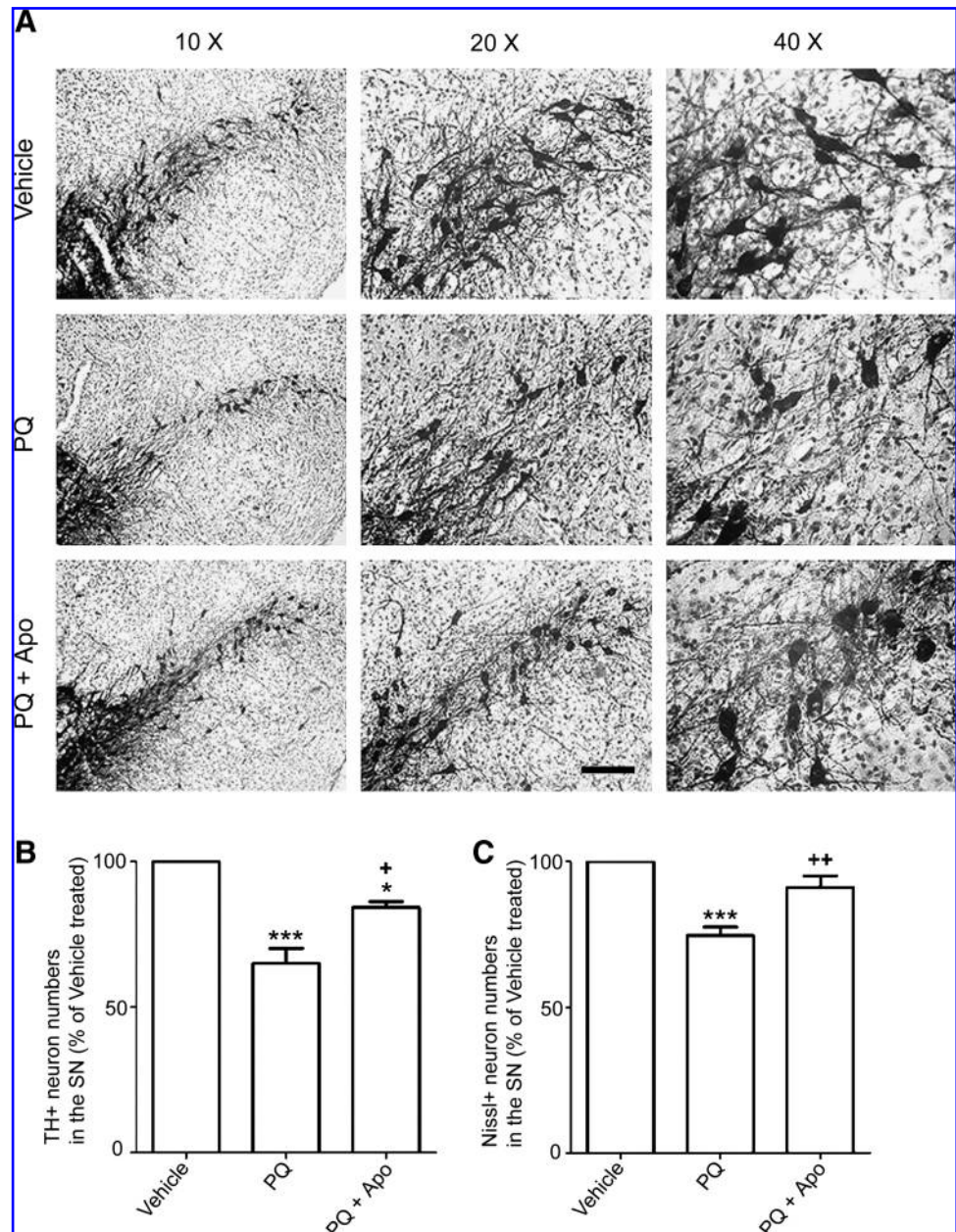
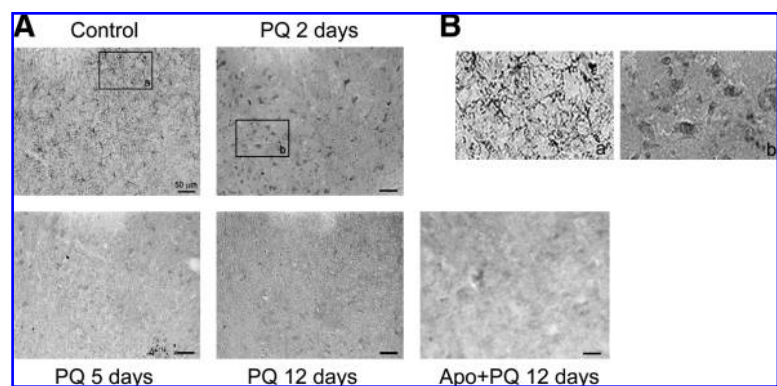


FIG. 8. The time-course decrease in CD11b-positive microglia in mice treated with PQ.

(A) Representative photomicrographs of four independent experiments depicting CD11b-positive microglial immunostaining in the SNpc of mice treated with vehicle, PQ, or Apo+PQ at days 2, 5, and 12. Changes are observed in the number of CD11b-immunoreactive cells in the SNpc after PQ and Apo+PQ treatment. (B) Higher magnification of respective boxed areas in (A). Control, vehicle treated; PQ 2, 5, 12 days, PQ treated; Apo+PQ 12 days, PQ and apocynin co-treatment. (Control, $n=5$; PQ group, $n=7$; Apo+PQ group, $n=5$). Scale bar, 50 μ m.



Microglia are another cellular component in brain parenchyma affected by PQ administration. PQ activates Nox-mediated superoxide generation in BV2 microglia (43). Primary neuron-microglia mixed culture treated with a low concentration of PQ (1 μ M) showed selective DA cell death, whereas microglia-depleted cultures or neuron-microglia mixed cultures of Nox2-null mice were resistant to PQ exposure. This result suggests that DA neurotoxicity is indirectly caused by low-dose PQ-mediated microglial Nox2 activation (59). In the present study, however, a significant decrease in the number of CD11b-positive microglia was observed in PQ-treated mice, suggesting that the PQ-injection paradigm used in the present study could be highly toxic to microglia.

It is well recognized that apocynin can act as a NADPH oxidase inhibitor, and its neuroprotective effects have been reported in several CNS disorders. ALS mice treated with apocynin showed an increase in their average life spans (27). In the ketamine-induced schizophrenia animal model, inhibition of NADPH oxidase by apocynin prevents the dysfunction of cortical inhibitory interneurons elicited by ketamine and reduced NADPH oxidase activity (8). In the experimental stroke model using middle cerebral artery occlusion, apocynin significantly prevents blood-brain barrier disruption (35). Recent studies, however, raised controversy that apocynin may work as an antioxidant instead of NADPH oxidase inhibitor, especially in the vascular system. The results suggest that it may have a tissue-specific role. Based on these facts, we first tested whether apocynin may reduce PQ-mediated ROS generation and cell death. Both PQ-induced ROS and DA cell death were significantly decreased by pretreatment with apocynin. We continued to investigate whether apocynin reduces SN DA neuronal death elicited by systemic administration of PQ. DA neuronal death in the SN was significantly reduced by apocynin. It was reported that in a mouse stroke model, the lower dose of apocynin administered IV works more effectively than the higher dose (56). It might be worth while to test several different doses of apocynin in a future study.

Next, we tested the specific role of Nox1 in the PQ-mediated ROS generation and DA cell death by siRNA-mediated Nox1 knockdown. Significant decreases in both ROS and cell death were achieved by Nox1 knockdown, suggesting that Nox1 plays a significant role in PQ-mediated oxidative stress and DA cell death. In addition to Nox1 in DA neurons, Nox2 in microglia may serve as a target for apocynin. Thus, the apocynin-mediated DA neuroprotection might be a combined outcome of microglial Nox2 and neuronal Nox1 inhibitions. To clarify the role of Nox1 in the nigrostriatal DA pathway degeneration caused by PQ, further study with Nox1-knockout mice or the *in vivo* knockdown system would be necessary.

In summary, the present study demonstrated that PQ may induce its toxic effects on DA neurons by activating the Nox system, particularly Nox1, which, in turn, generates ROS and eventually results in DA neuronal death. The study also raises the possibility that dopaminergic Nox1 may serve as a novel target for pharmacologic intervention of PD.

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Author Disclosure Statement

No competing financial interests exist.

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Abbreviations Used

Apo = apocynin
 CNS = central nervous system
 DA = dopaminergic
 DCFDA = 2',7'-dichlorodihydrofluorescein diacetate
 DHE = dihydroethidium
 DMSO = dimethyl sulfoxide
 FCCP = carbonylcyanide-4-(trifluoromethoxy)-phenylhydrazone
 GAPDH = glyceraldehyde 3-phosphate dehydrogenase
 IP = intraperitoneal
 LDH = lactate dehydrogenase
 MPP⁺ = 1-methyl-4-phenylpyridinium
 MPTP = 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine
 MTT = 3-(4,5-dimethylthiazal-2-yl)-2,5-diphenyl-tetrazolium bromide
 NADPH = nicotinamide adenine dinucleotide phosphate
 NBT = nitroblue tetrazolium
 NO = nitrogen oxide
 Nox = NADPH oxidase
 Nox1 = NADPH oxidase 1
 Nox2 = gp91^{phox}
 Nox3 = NADPH oxidase 3
 Nox4 = NADPH oxidase 4 OH⁻, hydroxyl radical
 PD = Parkinson's disease
 PQ = paraquat
 ROS = reactive oxygen species
 SN = substantia nigra
 SNpc = SN pars compacta
 TH = tyrosine hydroxylase

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