Abstract. Ischemia-reperfusion (I-R) injury is involved in the pathogenesis of several human diseases. In the present study, the kinetics of the H$_2$S producing enzymes-nuclear factor erythroid 2-like 2 (Nrf2)-antioxidant proteins axis under anoxia or reoxygenation was evaluated, as well as its effects on survival of mouse renal proximal tubular epithelial cells (RPTECs). In RPTECs subjected to anoxia and subsequent reoxygenation, reactive oxygen species (ROS) production, lipid peroxidation, ferroptotic cell death, the levels of the H$_2$S producing enzymes and H$_2$S, the expression of Nrf2 and its transcriptional targets superoxide dismutase-3, glutathione reductase, ferritin H and cystine-glutamate antiporter, as well as apoptosis, and the levels of p53, Bax and phosphorylated p53 were assessed. When needed, the H$_2$S producing enzyme inhibitor amino-oxyacetate, or the ferroptosis inhibitor α-tocopherol, were used. Reoxygenation induced ferroptosis, whereas anoxia activated the p53-Bax pathway and induced apoptosis. The H$_2$S producing enzymes-Nrf2-antioxidant proteins axis was activated only during anoxia and not during reoxygenation, when cellular viability is threatened by ROS overproduction and the ensuing ferroptosis. The activation of the above axis during anoxia ameliorated the effects of the apoptotic p53-Bax pathway, but did not adequately protect against apoptosis. In conclusion, the H$_2$S-Nrf2 axis is activated by anoxia, and although it reduces apoptosis, it does not completely prevent apoptotic cell death. Determining the exact molecular mechanisms involved in reoxygenation-induced cell death may assist in the development of clinically relevant interventions for preventing I-R injury.

Introduction

Ischemia-reperfusion (I-R) injury is implicated in several human diseases, and consists of two phases: ischemia and reperfusion. Occlusion of an artery or a decreased effective blood volume results in the ischemic phase. During the ischemic period, cell injury ensues as a result of anoxia-induced cellular energy collapse (1-3). Recanalization of the occluded artery or restoration of the effective blood volume restores blood supply leading to the reperfusion phase. During the reperfusion period, the re-entry of oxygen to the ischemic tissue results in a burst of reactive oxygen species (ROS) production and eventually in cell injury and death (1-3). Similar to other organs, such as the brain and the heart (1-3), the kidney is extremely vulnerable to I-R injury, with this type of injury being the leading cause of acute kidney injury (4).

The transcription factor nuclear factor erythroid 2-related factor 2 (Nrf2) regulates the transcription of several antioxidant and anti-ferroptotic genes (6,7), and it has been shown to protect against ferroptosis during reoxygenation. This is particularly true in RPTECs derived from the hibernator Syrian hamster (8,9). Hibernation involves cycles of torpor, characterized by a notable decline in heart and breathing rates, as well as in blood pressure, interrupted by interbout arousals and restoration of the above parameters. Hence, unlike most mammals, hibernators resist repeated cycles of I-R (10,11). In our previous study, it was shown that Nrf2 is activated during reoxygenation and is cytoprotective against ferroptosis in Syrian hamster RPTECs subjected to warm anoxia and subsequent reoxygenation (8). However, the mechanism underlying an insufficiency of the above system to protect non-hibernating mammals from reoxygenation-induced cell death remains to be determined.
Hydrogen sulfide (H$_2$S) serves a significant role in Nrf2 activation. H$_2$S is produced by the enzymes cystathionine β-synthase (CBS), cystathionine γ-lyase (CSE) and 3-mercaptopyruvate sulfurtransferase (3-MST) (12). H$_2$S interacts with, and induces conformational changes in Kelch-like ECH-associated protein 1 (Keap1), resulting in the release of Nrf2 from the Nrf2-Keap1 complex and rescuing Nrf2 from proteasomal degradation (13,14). Following release from Keap1, Nrf2 translocates to the nucleus and increases transcription of several antioxidant genes (6,7). This process is activated by reoxygenation in hibernating species offering protection against cell injury. It has been shown that all the aforementioned H$_2$S-producing enzymes, as well as the H$_2$S levels, are upregulated during reoxygenation and this results in the activation of Nrf2 in the context of resistance to reoxygenation-induced cell death in Syrian hamster RPTECs (8).

Mouse RPTECs die during anoxia through apoptosis (5). The role of the H2S-Nrf2 axis in apoptotic cell death is contradictory. Previous studies have shown that endogenous H$_2$S or Nrf2 activation protects against apoptosis (15-17); whereas other studies have shown that H$_2$S induces apoptosis (18-20). One of the latter studies which showed H$_2$S-induced apoptosis, suggesting that apoptosis was mediated by the activation of the proapoptotic p53/Bcl-2-associated X protein (Bax) pathway (20).

In support of the protective role of the H$_2$S-Nrf2-antioxidant proteins axis, numerous studies have shown that exogenous sulfide donors protect organs against I-R injury (21-24), including the kidneys (25-27).

To evaluate the possible role of the H$_2$S-Nrf2-antioxidant proteins axis in protecting non-hibernator mammals against I-R injury, the kinetics of the above axis in RPTECs derived from the non-hibernator mouse subjected to anoxia or reoxygenation were assessed. The non-specific inhibitor aminooxyacetate (AOAA) was used as a H$_2$S production inhibitor. AOAA directly inhibits CBS and CSE (28), and 3-MST indirectly, as 3-MST converts cysteine to pyruvate with the assistance of cysteine aminotransferase, and AOAA inhibits transamination (29). Additionally, when needed, the lipid peroxidation and ferroptosis inhibitor α-tocopherol was used (30).

Materials and methods

Cell culture and treatment. Primary C57BL/6 mouse RPTECs (cat. no. C57-6015, Cell Biologies, Inc.) were cultured in Complete Epithelial Cell Medium kit, supplemented with epithelial cell growth supplement (0.1% epithelial growth factor, Complete Epithelial Cell Medium kit, supplemented with (cat. no. C57-6015, Cell Biologics, Inc.) were cultured in a humidified atmosphere containing 5% CO$_2$ at 37°C for 2 h. To simulate reperfusion, after 24 h of anoxia, RPTECs were removed from the Anaerobe Container system and washed with PBS (Sigma-Aldrich; Merck KGaA), the culture medium was replaced with fresh medium, and the cells were cultured in a humidified atmosphere containing 5% CO$_2$ at 37°C for 2 h.

The periods of anoxia and reoxygenation were selected based on our previous study, in which it was shown that primary mouse RPTECs viability declines considerably after 48 h of anoxia and 4 h of reoxygenation (5). Cells were harvested after 24 h of anoxia or after 2 h of reoxygenation, as past these time points, the cell condition was deteriorated considerably, with the majority of cells not being suitable for further analysis (5). Each experiment was repeated six times.

For evaluating ferroptosis, 100 µM of α-tocopherol (Sigma-Aldrich; Merck KGaA) was added to inhibit lipid peroxidation and ferroptosis. For assessing the effect of H$_2$S, 2 mM AOAA (Selleck Chemicals) was used to inhibit H$_2$S-producing enzymes. The above AOAA concentration was selected after assessing its cytotoxicity in mouse RPTECs, as described below.

AOAA cytotoxicity in RPTECs. Mouse RPTECs were cultured in 96-well plates in a humidified atmosphere containing 5% CO$_2$ in the presence/absence of AOAA at a concentration of 0.5, 1 or 2 mM for 24 h. A lactate dehydrogenase (LDH) release assay was performed to assess cytotoxicity using a Cytoxtox Non-Radioactive Cytotoxicity assay kit (Promega Corporation). Cell necrosis was calculated using the following formula: Cell necrosis (%)=(LDH in the supernatant/total LDH) x100. Experiments were repeated six times.

Evaluation of proteins of interest. Mouse RPTECs were cultured in 6-well plates. Following anoxia and/or reoxygenation, RPTECs were lysed using T-PER tissue protein extraction reagent (Thermo Fisher Scientific Inc.) supplemented with protease (Sigma-Aldrich; Merck KGaA) and phosphatase inhibitors (Roche Diagnostics). After protein quantification using a Bradford assay (Sigma-Aldrich; Merck KGaA), 10 µg of protein from each sample was used for western blotting. Proteins were electrophoresed using a 4-12% bis-tris acrylamide gels (cat. no. NP0323BOX, NuPAGE 4-12% Bis-Tris Gel 1 mm x 15 well; Invitrogen; Thermo Fisher Scientific, Inc.). Proteins were transferred to a PVDF membrane, and membranes were blocked using skimmed milk in Tris-buffered saline with Tween-20. Blots were incubated with the primary antibody against the protein of interest for 16 h at 4°C, followed by incubation with the secondary antibody incubation for 30 min at room temperature. The LumiSensor Plus Chemiluminescent HRP Substrate kit (GenScript) was used for enhanced chemiluminescent detection of the bands. The Restore Western Blot Stripping Buffer kit (GenScript) was used whenever reprobing of the PVDF blots was required. Densitometry analysis was performed using ImageJ version 1.51t (National Institutes of Health). Experiments were repeated six times.

Primary antibodies used were specific for CBS (1:1,000; cat. no. TA338394; OriGene Technologies Inc.), CSE (1:100; cat. no. sc-374249; Santa Cruz Biotechnology, Inc.), 3-MST (1:100; cat. no. sc-376168; Santa Cruz Biotechnology, Inc.), Nrf2 (1:1,000; cat. no. TA343586; OriGene Technologies, Inc.),
superoxide dismutase 3 (SOD3; 1:100; cat. no. sc-271170; Santa Cruz Biotechnology, Inc.), glutathione reductase (GR; 1:100; cat. no. sc-133245; Santa Cruz Biotechnology, Inc.), ferritin heavy chain (1:100; cat. no. sc-376594; Santa Cruz Biotechnology, Inc.), cystine-glutamate antiporter (xCT; 1:1,000; cat. no. ANT-111; Alomone Labs), activated cleaved-caspase-3 (175) (1:500; cat. no. 7074; Cell Signaling Technology, Inc.), p53 (1:500; cat. no. 2524 Cell Signaling Technology, Inc.), p53 phosphorylated at serine 15 (p-p53) (1:500; cat. no. 9284; Cell Signaling Technology, Inc.), Bax (1:500; cat. no. 5023; Cell Signaling Technology, Inc.) and β-actin (1:2,500; cat. no. 4967; Cell Signaling Technology, Inc.). Horseradish peroxidase-conjugated anti-rabbit IgG (1:1,000; cat. no. 7074; Cell Signaling Technology, Inc.) or horseradish peroxidase-conjugated anti-mouse IgG (1:1,000, cat. no. 7076; Cell Signaling Technology, Inc.) were used as secondary antibodies.

Assessment of H$_2$S production. At the end of the 24-h anoxia period and the 2-h reoxygenation periods, H$_2$S production was assessed by measuring its concentration in the supernatants of RPTECs cultured in 6-well plates. The effect of the H$_2$S-producing enzymes inhibitor AOAA was also assessed. H$_2$S production was measured using a methylene blue assay as described previously (31,32). Zinc acetate (1% w/v) (Sigma-Aldrich; Merck KGaA) was added immediately to 1 ml of each supernatant to trap the produced H$_2$S. N,N-dimethyl-p-phenylenediamine dihydrochloride (400 mg) (Sigma-Aldrich; Merck KGaA) was dissolved in 10 ml 6 M HCl, ferric chloride (600 mg; Sigma-Aldrich; Merck KGaA) in 10 ml 6 M HCl, and then, 1 ml from each of the two solutions were mixed to prepare the required diamine-ferric solution. Subsequently, 50 µl diamine-ferric solution was added to each supernatant for 30 min and incubated at 37°C, 200 µl of each reaction was placed in the wells of a 96-well plate, and the amount of methylene blue formed in each supernatant was measured at 670 nm on an EnSpire® Multimode Plate Reader (PerkinElmer, Inc.). To extrapolate the results of each experimental reaction, different concentrations of methylene blue were also measured to create a standard curve (Merck KGaA). These experiments were repeated six times.

Evaluation of ROS production. ROS production was measured in RPTECs cultured in 96-well plates. Following anoxia and reoxygenation, 5 µM CellROX® Deep Red Reagent (Invitrogen; Thermo Fisher Scientific, Inc.), a fluorogenic probe, was added to the culture medium and cells were incubated at 37°C for 30 min. Subsequently, RPTECs were washed with PBS, and an EnSpire® Multimode Plate Reader was used to measure fluorescence signal intensity. These experiments were repeated six times.

Assessment of lipid peroxidation. Lipid peroxidation was assessed in RPTECs cultured in 6-well-plates. Malondialdehyde (MDA), the end-product of lipid peroxidation, was measured fluorometrically in cell extracts with a Lipid Peroxidation (MDA) assay kit (cat. no. abl18970; Abcam). The kit fluorometrically detects MDA levels as low as 0.1 nmol. A Bradford assay was performed prior to MDA measurement, and the lysate volumes of all samples were adjusted to a protein concentration of 1 mg/ml. These experiments were repeated six times.

Assessment of cell ferroptosis. In mouse RPTECs subjected to reoxygenation, cell death ensues via ferroptosis. To assess ferroptosis, at the end of the reoxygenation period, cell necrosis was assessed in RPTECs cultured in 96-well plates, in the presence/absence of lipid peroxidation and ferroptosis inhibitor α-tocopherol. The effects of the H$_2$S-producing enzymes inhibitor AOAA was also evaluated. An LDH release assay was performed for assessing cell necrosis using the Cytotox Non-Radioactive Cytotoxic assay kit. Cell necrosis was calculated as follows: Cell necrosis (%) = (LDH in the supernatant/total LDH)x100. These experiments were repeated six times.

Statistical analysis. SPSS Version 20 (IBM, Corp.) was used for statistical analysis. A one-sample Kolmogorov-Smirnov test was used to confirm that the evaluated variables were normally distributed. For comparison of means, a one-way ANOVA followed by Bonferroni's correction test was used. Results are expressed as the mean ± standard error of mean. P<0.05 was considered to indicate a statistically significant difference.

Results

AOAA is not toxic for RPTECs at the assessed concentrations. AOAA did not exhibit any notable cytotoxic effects at any of the tested concentrations (Fig. 1). A concentration of 2 mM was selected for all subsequent experiments.

ROS production, lipid peroxidation, ferroptosis and the role of endogenous H$_2$S. Compared with the control RPTECs, in RPTECs cultured under anoxic conditions, ROS production did not differ significantly (signal intensity 40.4±0.4 vs. 39.6±1.5, respectively). However, reoxygenation increased ROS production significantly (signal intensity 71.0±3.2; P<0.001 compared with control and anoxia; Fig. 2A).

Anoxia did not significantly affect lipid peroxidation, whereas reoxygenation induced lipid peroxidation in RPTECs. MDA levels were 4.0±0.1 µM in control cells and 4.0±0.1 µM in RPTECs cultured under anoxic conditions. Reoxygenation increased MDA levels significantly to 51.7±1.2 µM (P<0.001 compared with control and anoxia; Fig. 2B).

In RPTECs cultured under normoxic conditions, the ferroptosis inhibitor α-tocopherol, the H$_2$S-producing enzymes inhibitor AOAA, or their combination did not significantly alter cell necrosis, which was 9.8±0.2, 9.6±0.3, 10.3±0.6 and 9.9±0.4%, respectively. Reoxygenation increased cell necrosis to 28.8±0.7% (P<0.001 compared with control; Fig. 2C).

Treatment of RPTECs subjected to reoxygenation with α-tocopherol ameliorated cell necrosis considerably (17.4±0.4%; P<0.001 compared with reoxygenation alone), whereas treatment with AOAA significantly increased cell necrosis (37.8±0.9%; P<0.001 compared with reoxygenation alone; Fig. 2C).

Collectively, these results show that reoxygenation increases ROS production and induces ferroptosis, and also suggests a protective role of endogenous H$_2$S against ferroptosis.
Kinetics of H₂S-producing enzymes expression and H₂S under anoxia and reoxygenation. The expression of all the H₂S-producing enzymes increased significantly under anoxia (P<0.001 compared with the control) and returned to baseline during reoxygenation. For CBS, the mean optical density (OD) was 5.4±0.9 under normoxia, 14.0±2.2 under anoxia and 5.6±0.7 under reoxygenation (Fig. 3A). The OD values of CSE were 7.6±0.5, 10.0±0.6 and 7.4±0.7 under normoxia, anoxia and reoxygenation, respectively (Fig. 3B). For 3-MST, the ODs were 7.1±1.0, 10.2±0.9 and 7.7±0.8, under normoxia, anoxia and reoxygenation, respectively (Fig. 3C).

Production of H₂S follows the pattern of H₂S-producing enzymes expression. Under control conditions, H₂S concentration was 9.5±0.3 µM, under anoxia it increased to 24.3±1.1 µM (P<0.001 compared with the control), and reoxygenation decreased H₂S concentration to 9.3±0.3 µM (P<0.05 compared with the control). The presence of AOAA reduced the H₂S concentration significantly under all conditions. In control cells, AOAA decreased H₂S concentration to 5.6±0.4 µM (P<0.001 compared with the control alone), in RPTECs under anoxia to 5.4±0.6 mM (P<0.001 compared with anoxia alone), and in RPTECs subjected to reoxygenation to 5.7±0.4 µM (P<0.001 compared with reoxygenation alone; Fig. 3D).

Hence, in RPTECs, all H₂S-producing enzymes, and thus H₂S production, were upregulated during anoxia and returned to baseline during reoxygenation. AOAA decreased H₂S production under all cell culture conditions.

Activation status of Nrf2-antioxidant proteins axis during anoxia and reoxygenation. The Nrf2 levels followed the fluctuations of H₂S concentrations under anoxia and reoxygenation. Nrf2 mean OD was 5.9±0.2 under control conditions, which increased to 14.4±1.5 under anoxia (P<0.001 compared with the control), and returned to baseline under reoxygenation (4.8±0.1; P>0.05 compared with the control; Fig. 4A).

The expression of all the evaluated antioxidant proteins, which are transcriptional targets of Nrf2, followed the fluctuations of Nrf2 levels. SOD-3 OD was 5.7±0.8 under control conditions, which increased to 13.0±0.5 under anoxia (P<0.001 compared with the control), and returned to baseline under reoxygenation (6.3±0.9; P<0.05 compared with the control; Fig. 4B). OD values for GR were 5.8±0.8 under normal conditions, 12.0±1.1 under anoxia (P<0.001 compared with the control) and 7.2±1.3 under reoxygenation (P<0.05 compared with the control; Fig. 4C). For Ferritin H the OD values were 6.1±0.5 under normal conditions, 12.6±0.4 under anoxia (P<0.001 compared with the control) and 6.3±0.7 under reoxygenation (P<0.05 compared with the control; Fig. 4D). Finally, for xCT the OD values were 6.3±1.4 under normal conditions, 12.9±1.8 under anoxia (P<0.001 compared with the control) and 5.8±1.3 under reoxygenation (P<0.05 compared with the control; Fig. 4E).

Therefore, in RPTECs, the anoxia-induced increase in H₂S production enhanced both Nrf2 levels and the expression of various antioxidant proteins, which is under the transcriptional control of Nrf2. During reoxygenation, when ROS levels increase and when the antioxidant proteins are required, the axis remained inactive.

Apoptosis during anoxia and reoxygenation. Apoptotic cell death was assessed by measuring the levels of activated cleaved-caspase-3, in which all the apoptotic pathways converge (33). Anoxia induced apoptosis in RPTECs, whereas reoxygenation did not. Cleaved-caspase-3 mean OD values 4.3±0.3 under control conditions, increased to 12.9±1.3 under anoxia (P<0.001 compared with the control) and returned to baseline levels (5.2±0.7; P>0.05 compared with the control) in RPTECs subjected to reoxygenation (Fig. 5).

Thus, in mouse RPTECs, apoptosis ensues only under anoxic conditions and not during the reoxygeanation phase.

Effect of H₂S on anoxia-induced apoptosis and on the pro-apoptotic p53-Bax axis. Since anoxia induces apoptosis in RPTECs, and various studies have produced contradictory results regarding the effects of endogenous H₂S on apoptosis, and on the proapoptotic p53-Bax axis, the impact of the H₂S-producing enzymes inhibitor AOAA on the above parameters in RPTECs subjected to anoxia were assessed.

In RPTECs subjected to anoxia, AOAA increased apoptosis further. The mean OD values of the cleaved-caspase-3 was 4.7±0.3 in the control RPTECs, which increased to 9.3±0.2 in RPTECs subjected to anoxia (P<0.001 compared with the control), and increased even further to 11.5±0.4 in RPTECs subjected to anoxia and treated with AOAA (P<0.001 compared with the control and anoxia). Additionally, under control conditions, AOAA enhanced cleaved-caspase-3 levels to 7.9±0.3 (P<0.001 compared with the control; Fig. 6A).

The increase in apoptosis due to AOAA treatment may result from the activation of the proapoptotic p53-Bax axis. The p53 OD values were 4.6±0.9 in control RPTECs, increased to 9.7±0.5 under anoxia (P<0.001 compared with the control), and further increased to 11.8±0.3 in the presence of AOAA (P<0.001 compared with the control and anoxia). Additionally, AOAA enhanced p53 levels to 9.7±0.3 under control conditions (P<0.001 compared with the control; Fig. 6B).

These alterations in p53 levels were likely the result of an increase in its phosphorylation, which dissociates p53 from mouse double minute 2 homolog (MDM2) and protects p53 from degradation. The increase in apoptosis due to AOAA treatment may result from the activation of the proapoptotic p53-Bax axis. The p53 OD values were 4.6±0.9 in control RPTECs, increased to 9.7±0.5 under anoxia (P<0.001 compared with the control), and further increased to 11.8±0.3 in the presence of AOAA (P<0.001 compared with the control and anoxia). Additionally, AOAA enhanced p53 levels to 9.7±0.3 under control conditions (P<0.001 compared with the control; Fig. 6B).

These alterations in p53 levels were likely the result of an increase in its phosphorylation, which dissociates p53 from mouse double minute 2 homolog (MDM2) and protects p53 from degradation. The increase in apoptosis due to AOAA treatment may result from the activation of the proapoptotic p53-Bax axis. The p53 OD values were 4.6±0.9 in control RPTECs, increased to 9.7±0.5 under anoxia (P<0.001 compared with the control), and further increased to 11.8±0.3 in the presence of AOAA (P<0.001 compared with the control and anoxia). Additionally, AOAA enhanced p53 levels to 9.7±0.3 under control conditions (P<0.001 compared with the control; Fig. 6B).
from proteasomal degradation. The p-p53 OD was 4.2±0.4 in control RPTECs, which increased to 8.9±0.2 under anoxia (P<0.001 compared with the control), and further to 13.7±0.7 when treated with AOAA (P<0.001 compared with the control and anoxia). AOAA treatment increased the p-p53 OD to 6.6±1.0 in RPTECs cultured under control conditions (P<0.001 compared with the control; Fig. 6C).

The expression of the proapoptotic protein Bax followed the pattern of p53, which controls Bax transcription. Anoxia increased Bax OD values from 2.4±0.5 in the control to 9.9±1.1 under anoxia (P<0.001), and Bax combined with AOAA treatment further increased Bax levels (15.5±0.7; P<0.001 compared with the control and anoxia). Treatment of RPTECs with AOAA under normoxia also increased Bax expression levels (5.5±0.5; P<0.001 compared with the control; Fig. 6D).

Based on these results, it is hypothesized that endogenous H2S, which increases under anoxic conditions, ameliorates anoxia-induced apoptosis by interfering with the p53-Bax axis. However, as the anoxia-induced changes proceed, apoptosis occurs indicating that the upregulation in H2S-producing enzymes during anoxia is insufficient for preventing apoptosis completely.

**Discussion**

In mouse RPTECs, ROS production was increased only during reoxygenation, when oxygen was available, resulting in lipid peroxidation and cell necrosis. The inhibitor of lipid peroxidation and ferroptosis, α-tocopherol, protected mouse RPTECs against reoxygenation-induced cell necrosis, indicating that during this phase, cell death occurs via ferroptosis (34). Similar results were obtained in a previous study with isolated mouse renal tubules subjected to anoxia and subsequent reoxygenation (35). Treatment with AOAA increased ferroptosis, underlying the protective role of H2S.
Expression of all the assessed H$_2$S producing enzymes, as well as H$_2$S levels, were upregulated during anoxia; the increased H$_2$S levels upregulated Nrf2 expression levels, and thus indirectly, the expression of the assessed transcriptional targets of Nrf2 (6,7,13,14). The targets assessed were SOD-3, which catalyzes the dismutation of superoxide radicals into either molecular oxygen or hydrogen peroxide (36); GR, which catalyzes the reduction of glutathione disulfide to the sulfhydryl form glutathione (GSH) (36); ferritin, which functions by sequestering intracellular labile iron preventing the Fenton reaction and blocking the function of iron-containing lipoxygenases (30); and the cystine-glutamate antiporter xCT, which is responsible for the entry of the cystine required for GSH synthesis into the cell and prevents ferroptosis (30). All the above Nrf-2 targets were upregulated during anoxia.

However, during reoxygenation, when the activation of the antioxidant defense system is required due to ROS overproduction, the H$_2$S producing enzymes-Nrf2-antioxidant proteins axis was downregulated to baseline levels, leaving the cells unprotected. Thus, during anoxia-reoxygenation injury, this mistimed activation of the above axis is unable to protect RPTECs against reoxygenation-induced ferroptotic cell death.

The significance of the above axis in protecting cells against reoxygenation-induced ferroptosis has been detected in hibernating species, which are subjected to repeated cycles of I-R during hibernation without experiencing any injuries (10,11). Contrary to mouse RPTECs, in Syrian hamster RPTECs, the activation of H$_2$S producing enzymes-Nrf2-antioxidant proteins axis takes place later, at the appropriate time during reoxygenation and ROS overproduction, protecting cells from ferroptotic cell death (8).

Furthermore, the results of the present study support the possible protective role of exogenous sulfur donors in the prevention of reoxygenation-induced cell death. Several studies in non-hibernating species have demonstrated the protective effects of such an intervention in experimental models of acute kidney injury (25-27), myocardial infarction (21), cerebral stroke (22) and multiorgan failure (23,24). Since oxidative stress is implicated in several renal pathologies, Nrf2 activators other than H$_2$S donors have also been assessed in experimental models of various kidney diseases with promising results, and several clinical studies are being performed (40). In an experimental model of I-R-induced kidney injury, the Nrf2 activator bardoxolone methyl ameliorated structural injury and renal dysfunction in mice (41). Bardoxolone imidazolide was also found to improve survival, renal function, kidney histology and production of pro-inflammatory cytokines in mice subjected to kidney I-R injury (42). Similar beneficial results were observed in mice administered omaveloxolone, another Nrf2 activator (43). Also, the interest in various natural products able to activate Nrf2 is growing. For example, the Nrf2 activator bardoxolone methyl ameliorated structural injury and renal dysfunction in mice (41). Bardoxolone imidazolide was also found to improve survival, renal function, kidney histology and production of pro-inflammatory cytokines in mice subjected to kidney I-R injury (42). Similar beneficial results were observed in mice administered omaveloxolone, another Nrf2 activator (43).
activator curcumin, was beneficial in a rat model of kidney I-R injury (44), and total flavonoids from Rosa laevigata Michx fruit upregulated Nrf2 expression, and exhibited beneficial effects in mice subjected in kidney I-R injury (45).

The results of the present study suggested that apoptosis, based on the levels of cleaved-caspase-3 in which all the apoptotic pathways converge (33), was only observed during anoxia, and not during reoxygenation. This is in agreement with the results of a previous study in which mouse RPTECs were subjected to similar experimental conditions (5).

Interestingly, in RPTECs under anoxia, administration of AOAA decreased H2S production, and in parallel, aggravated apoptotic cell death.

Under anoxia, in RPTECs, the levels of cleaved-caspase-3, p53 and Bax also increased, suggesting that apoptosis may be mediated by the p53-Bax pro-apoptotic pathway (46). The role of the p53-Bax pathway in I-R injury has been established, and in a previous study, transient silencing of p53 with a specific siRNA, protected rat kidney function from I-R injury by preventing apoptosis of RPTECs (47).

The results of two ongoing related clinical trials using a p53-specific siRNA; a phase 2 clinical trial on preventing acute kidney injury following cardiac surgery (clinicaltrials.gov/ct2/show/results/NCT02610283), and a phase 3 clinical trial on preventing delayed graft function following kidney transplantation from old donors (clinicaltrials.gov/ct2/show/results/NCT02610296) will clarify the clinical significance of p53 in I-R-induced apoptosis.

The levels of p-p53 were also increased during anoxia, suggesting that the increase in p53 levels was the result of p53 dissociation from the ubiquitin ligase MDM2 (46). The enhanced phosphorylation of p53 may result from anoxia-induced DNA damage, and the activation of a group of kinases which are implicated in the genome integrity checkpoint (46).

AOAA treatment resulted in an increase in the levels of p-p53, p53, Bax and CC3, suggesting a protective role of the H2S-Nrf2 axis against apoptosis. Interestingly, there is a

Figure 4. Expression of Nrf2 and its transcriptional targets under anoxia or reoxygenation. (A) Expression of Nrf2 and its transcriptional targets (B) SOD-3, (C) glutathione reductase, (D) ferritin H and (E) xCT increased under anoxia, and returned to baseline levels during reoxygenation. Representative blots are presented. *P<0.001 vs. Ctrl; †P<0.001 vs. Reox. Nrf2, nuclear factor erythroid 2-like 2; Ctrl, control; Reox, reoxygenation; OD, optical density; SOD-3, superoxide dismutase 3; xCT, cystine-glutamate antiporter.
The bidirectional interaction between Nrf2 and the p53 pathway. Nrf2 transcribes MDM2, which acts as a specific ubiquitin ligase and downregulates p53. p53 directly suppresses Nrf2, although p53 also transcribes p21, which is known to upregulate Nrf2 (48). The exact molecular interactions that transpire during anoxia remains to be elucidated.

The results of the present study are in agreement with studies showing an anti-apoptotic role of the H2S-Nrf2 axis (15-17), contradicting studies which showed the opposite outcomes (18-20). However, by interpreting the outcome of the experiments, which indicate that despite anoxia-induced activation of the H2S-Nrf2 axis, apoptosis occurs, it is likely that this system by itself is unable to confer full protection on RPTECs against anoxia-induced apoptosis. The latter highlights the possibility of exogenous sulfur donors to protect RPTECs against anoxia-induced apoptotic cell death. Combinations of sulfur donors with apoptosis inhibitors may prove more effective in the prevention or amelioration of I-R injury. Interestingly, and contrary to the mouse RPTECs, in RPTECs derived from the hibernator Syrian hamster, which resists apoptosis, the H2S-Nrf2 axis is activated at a later stage, during reoxygenation, suggesting that in hibernating species, the above axis is not responsible for the evident resistance to anoxia-induced apoptosis (8,9).
A limitation of the present study is the in vitro nature of the experiments. However, the strictly controlled experimental conditions allowed the study of the two different, subsequent, but distinct components of I-R injury separately, and to assess the different kinetics of the H$_2$S producing enzymes-Nrf2-antioxidant proteins axis under anoxia and reoxygenation, as well as its effect on cell survival. Thus, our results may be considered a starting point for further studies on the molecular mechanisms that govern the activity of the above axis under anoxia and reoxygenation, as well as for interventional in vivo studies.

In conclusion, the results of the present study suggest that in RPTECs, the H$_2$S-Nrf2 axis is activated by anoxia, and although it ameliorates apoptosis, it does not completely prevent apoptotic cell death, and is eventually overwhelmed. On the contrary, under reoxygenation, when the sudden increase in ROS production occurs, the antioxidant defense is essential for the protection of cells against ferroptotic cell death, the H$_2$S producing enzymes-Nrf2-antioxidant proteins axis is not upregulated. This mistimed activation of the above axis contributes to reoxygenation-induced cell death. Clarifying the precise molecular mechanisms underlying the mistimed H$_2$S producing enzymes-Nrf2-antioxidant proteins axis activation may result in clinically useful interventions for preventing I-R injury.

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Availability of data and materials

The datasets used and/or analyzed during the present study are available from the corresponding author on reasonable request.

Authors' contribution

TE designed the study, GP and TE performed the experiments. TE, GP, EN, GF, VL and IS analyzed the results. TE and GP wrote the manuscript. All authors approved the final manuscript.

Ethics approval and consent to participate

Not applicable.

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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