Exogenous hydrogen sulfide protects against hepatic ischemia/reperfusion injury by inhibiting endoplasmic reticulum stress and cell apoptosis

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Abstract. The aim of the present study was to explore the effect of exogenous hydrogen sulphide (H₂S) on endoplasmic reticulum (ER) stress (ERS) in a rat model of hepatic ischemia/reperfusion (I/R) injury. A total of 48 Sprague-Dawley rats were randomly divided into four groups (n=12/group) as follows: Sham, I/R, I/R preceded by NaHS (I/R-NaHS) and I/R preceded by L-C-propargylglycine (PAG), a H₂S inhibitor (I/R-PAG). With the exception of the sham group, the rats in the other groups were subjected to 30 min hepatic warm ischemia followed by reperfusion for 6 or 12 h. Hepatic function was evaluated by serum concentrations of alanine aminotransferase (ALT). Apoptosis of hepatic cells was assessed by TUNEL staining and measurement of caspase-12 expression. The expression levels of ERS-associated proteins and mRNAs of pancreatic ER eukaryotic translation initiation factor-2a kinase (PERK), activating transcription factor-6 (ATF6), glucose-regulated protein (GRP) 78, TNF-receptor-associated factor (TRAF)-2, C/EBP homologous protein (CHOP) and caspase-12 were also measured by western blotting and reverse transcription-quantitative PCR. The serum concentrations of ALT in the I/R and I/R-PAG groups were found to be significantly higher compared with those in the sham and I/R-NaHS groups after 6 h of reperfusion; in addition, the ALT level returned to normal in the I/R group, while it increased further in the I/R-PAG group after 12 h of reperfusion. A higher cell apoptosis rate was observed in the I/R and I/R-PAG groups and the highest cell apoptosis rate was observed in the I/R-PAG group; correspondingly, the expression of caspase-12 was increased in the I/R and I/R-PAG groups. H₂S appeared to significantly attenuate hepatic I/R-induced ERS response, as indicated by the decreased expression of ATF6, PERK, GRP78, TRAF2 and CHOP. Endogenous H₂S may serve a hepatoprotective function after I/R, and inhibition of endogenous H₂S results in aggravation of I/R damage. Exogenous H₂S was shown to inhibit ERS-related gene expression, leading to suppression of inflammatory reaction and improvement of I/R damage. Therefore, exogenous H₂S has therapeutic potential to alleviate hepatic I/R injury.

Introduction

Liver injury after the onset of reperfusion is the result of the interplay between different complex mechanisms, with cellular damage occurring during both the ischemic and reperfusion phases. Hepatic ischemia/reperfusion (I/R) causes major injury following vascular occlusion during both liver surgery and liver transplantation (1); however, its pathogenesis is largely unknown. The destructive effects of I/R may be triggered by the acute generation of reactive oxygen species (ROS) and nitrogen species after reoxygenation, which cause a chain reaction of cellular responses leading to inflammation, cell death and, eventually, organ failure (2-4). Several mediators, such as proinflammatory cytokines, ROS, adhesion molecules, chemokines and excess nitric oxide (NO) contribute to this injury (5). However, it was previously demonstrated that reperfusion did not induce oxidative stress, but sustained endoplasmic reticulum (ER) stress (ERS) in the livers of rats subjected to traumatic-hemorrhagic shock (6). The reason may be that the model used in that study was produced by bleeding until decompensation, followed by an inadequate or adequate reperfusion phase.

The ER is an organelle with important signaling and homeostatic functions, which is responsible for protein
folding, maturation and trafficking. ERS is a particular subcellular pathological process involving an imbalance of homeostasis and ER disorder. In the early stages of ERS, cells exhibit a protective unfolded protein response that changes the cellular transcriptional and translational programs to alleviate this process (7). ERS triggers an adaptive response pathway, named the unfolded protein response (UPR) pathway, through activation of pancreatic ER eukaryotic translation initiation factor-2α kinase (PERK), inositol requiring enzyme-1 (IRE1) and activating transcription factor-6 (ATF6). Several conditions, including hypoxia, I/R injury, neurodegeneration, heart disease and diabetes, may be associated with activation of ER-initiated cell death pathways (8). In samples from human livers subjected to I/R, UPR pathway activation was observed (9). Therefore, ERS may play an important role in hepatic I/R injury.

Endogenous hydrogen sulfide (H₂S) is increasingly being recognized as an important gaseous physiological mediator synthesized from cysteine by cystathionine γ lyase (CSE) and other naturally occurring enzymes. CSE is differentially expressed in the cardiovascular, neuronal, immune, renal, respiratory, gastrointestinal, reproductive, hepatic and endocrine systems (10). H₂S serves important functions in cytoprotection, anti-inflammation, nociceptive stimulation, regulation of insulin release and longevity (11‑13). Acute administration of H₂S, either prior to ischemia or at reperfusion, ameliorates hepatic I/R injury through an upregulation of intracellular antioxidant and antiapoptotic signaling pathways (14). However, whether ERS is involved in hepatic I/R injury remains unclear.

It was hypothesized that exogenous/endogenous H₂S inhibited ERS through limiting the extent of I/R injury. The present study was undertaken to investigate the effects of exogenous H₂S on I/R injury using an in vivo rat model. The aim was to determine whether exogenous H₂S can inhibit the expression of ERS- and cell apoptosis‑related genes, thereby leading to suppression of inflammatory reactions and improvement of I/R damage, and whether inhibition of endogenous H₂S may result in aggravation of I/R injury.

Materials and methods

Animal model. In total, 48 Sprague-Dawley male rats, aged 10 weeks (weighing 225-275 g) were supplied by The Animal Centre of Xining, Qinghai Province and housed in groups of two to four in 40x60x30 cm plastic cages with soft bedding under a 12‑h day and night cycle. The room temperature was maintained at 22±1°C and relative humidity was 45‑60%; water and food pellets were available ad libitum. Animal studies were performed in compliance with the guidelines of animal ethics of Qinghai University and were approved by the Ethics Committee Qinghai University Affiliated Hospital (approval no. 20180152; Xining, China).

Experimental design and surgery. A total of 48 Sprague-Dawley rats were randomly divided into four groups (n=12 group) as follows: Sham, I/R, I/R preceded by NaHS (I/R-NaHS) and I/R preceded by L-C-propargylglycine (PAG), a H₂S inhibitor (I/R-PAG). All surgeries were performed as previously described (15). All animals were anesthetized with 5% isoflurane for induction and 2-3% isoflurane for maintenance in an acrylic anesthesia chamber, and the liver was exposed through a transverse abdominal incision. A 30‑min period of hepatic warm ischemia was introduced by cross-clamping the hepatoduodenal ligament with a microvascular clamp (Pringle's maneuver) in the rats, with the exception of those in the sham group. Subsequently, the clamps were removed to allow reperfusion for 6 or 12 h before sampling, in order to explore whether the I/R injury could be reversed under the protective effect of H₂S. The animals of the I/R group were administered saline (1 ml/kg body weight) intraperitoneally before closing the incision. The animals of the I/R-NaHS group were administered a single dose (56 µmol/kg) of NaHS intraperitoneally 30 min before closing the incision (the dosage was set according to our preliminary experiment). To completely inhibit endogenous H₂S, the animals of the I/R-PAG group were administered 50 mg/kg of PAG intraperitoneally 30 min before and after closure of the incision. After the experiments were completed, all the rats were euthanized with 8% isoflurane inhalation for 30 min until breathing had stopped and their skin became cyanotic.

Serum biochemical analysis. At the end of the reperfusion period, 100-µl blood samples were collected from the aorta under anesthesia using heparinized syringes. The rats were euthanized after blood sample collection and liver samples were harvested. After 300 x g centrifugation at 4°C for 30 min, plasma was separated and stored at -20°C until analysis. Serum concentrations of alanine aminotransferase (ALT; cat. no. E-BC-K235-M, Elabscience Biotechnology, Inc.) were measured at 37°C using standard enzymatic techniques (Ektachem 700 analyzer; Kodak).

TUNEL assay. The harvested liver samples were snap-frozen in liquid nitrogen and stored in -80°C. Subsequently, the frozen liver samples were cut into 5-µm sections, and in situ apoptosis was detected using a TUNEL assay kit (Promega Corporation): Proteinase K was used to incubate the liver tissue samples for 4°C 20 min, before 50 µl terminal deoxynucleotidyl transferase incubation buffer was utilized to incubate the samples for 1 h at 37°C in the dark after the samples were washed for three times with PBS, 0.5 µg/ml DAPI (Beyotime Institute of Biotechnology) was used to stain the nucleus for 15 min at 4°C. The slides were dehydrated in 100% ethanol, cleared with xylene and mounted with 50 µl DAPI (Beyotime Institute of Biotechnology). Cells displaying brown staining within the nucleus were counted as apoptotic. The number of apoptotic cells was counted by an investigator who was blinded to the group assignment by examining three non-overlapping microscopic fields of view under high-power magnification (x400) via a Zeiss confocal Axioskop 2 Plus microscope (Carl Zeiss AG) and expressed as percentage (16).

Western blot analysis. Ischemic hepatic lobes were dissected and homogenized in a buffer [2% SDS, 50 mM DTT, 62.5 mM Tris (pH 6.8), 10% glycerol] containing 1 mM PMSF (Amresco, LLC). The samples were then boiled for 5 min and centrifuged at 7,500 x g for 10 min at 4°C. Supernatants
were collected and protein concentrations were determined using a Bradford Protein Assay kit (Beyotime Institute of Biotechnology). Protein samples (35 µg) were then loaded and separated on a 12% SDS-PAGE. Following transfer onto a nitrocellulose membrane, the membranes were incubated with 5% skimmed milk to block non-specific antigens at room temperature for 2 h and probed with rabbit polyclonal antibodies (all 1:1,000) against ATF6 (cat. no. ab203119, Abcam), PERK (cat. no. ab65142, Abcam), glucose-regulated protein 78 (ab21685, Abcam), TNF-receptor-associated factor 2 (TRAF)-2 (cat. no. ab62488, Abcam), C/EBP homologous protein (CHOP; cat. no. ab11419, Abcam), caspase-12 (cat. no. ab62463, Abcam) and GAPDH (cat. no. ab9485, Abcam) overnight at 4˚C. Following incubation with 1:1,000 diluted goat anti-rabbit IgG (H&L) secondary antibody (cat. no. G-21234, Thermo Fisher Scientific) for 2 h at room temperature, the protein bands were developed using an enhanced chemiluminescence kit (Amersham; Cytiva).

Reverse transcription-quantitative (RT-q) PCR analysis. Total RNA was isolated from liver tissue using TRIzol® reagent (Thermo Fisher Scientific, Inc.) according to the manufacturer's instructions and reversely transcribed using PrimeScript™ RT reagent kit (Takara Bio, Inc.) at 37˚C for 45 min and then 95˚C for 5 min. cDNA was used as the template for the qPCR reaction. The PCR conditions were as follows: Step 1, pre-denaturing at 95˚C for 30 sec; step 2, 40 cycles at 94˚C for 5 sec, followed by 60˚C for 30 sec. qPCR was performed with SYBR™ Green Master Mix kit (cat. no. 4344463; Thermo Fisher Scientific, Inc.) using AB 2900HT fluorescence ration PCR instrument. The primers for qPCR amplification are listed in Table I. For determination of relative quantification, the expression levels of the genes were calculated and expressed based on the 2\(^{-\Delta\Delta Cq}\) method (17). GAPDH was selected as an internal control gene.

Statistical analysis. Data are expressed as mean ± SEM of three separate repeated experiments. Two-way ANOVA followed by Bonferroni's post hoc test was performed to assess variation among experimental groups. P<0.05 was considered to indicate statistically significant differences. All analyses were performed with SPSS 16.0 (SPSS Inc.).

Results

\(H₂S\) restores liver function following I/R and inhibition of endogenous \(H₂S\) results in aggravation of I/R damage. In order to assess hepatocellular damage due to I/R, the serum ALT concentration was measured. A marked increase in the

Table I. Primers for quantitative PCR amplification.

| Gene      | Primer sequence (5’→3’)       | Expected size (bp) | GenBank accession no. |
|-----------|-------------------------------|--------------------|-----------------------|
| ATF6      | Forward: GGCTTCCTCAGGTTGTCCTCC  | 177                | NM_001107196.1        |
|           | Reverse: GTCAGGCTGTCCCTCTCC    |                    |                       |
| PERK      | Forward: TGGAGGAGGAGAGGAAATG   | 215                | NM_031599.2           |
|           | Reverse: GAGAGCAGAAGGAGGTTATGG |                    |                       |
| GRP78     | Forward: TAATCGCCACCGGATATG    | 193                | NM_013083.2           |
|           | Reverse: TTGTCAGTCGAGGTATGG    |                    |                       |
| TRAF2     | Forward: TGATGCTTCAGGAGGATG    | 101                | NM_01107815.2         |
|           | Reverse: TGCTTCGGTTGTTATGG     |                    |                       |
| CHOP      | Forward: GGAGAAAGGCAGGGAGAATG  | 176                | NM_01109986.1         |
|           | Reverse: GAGACACAGAAGGAGTTATGG |                    |                       |
| Caspase-12| Forward: ACTGTCCGACTCGAGAAC     | 104                | NM_130422.1           |
|           | Reverse: AGTGGCTATCCCCTTGTG    |                    |                       |
| GAPDH     | Forward: TGCCACTCAGAAGACTGGG   | 85                 | AF106860.2            |
|           | Reverse: GAGATGAGAGATGTTTCT   |                    |                       |

PERK, pancreatic ER eukaryotic translation initiation factor-2a kinase; ATF6, activating transcription factor-6; GRP78, glucose-regulated protein 78; TRAF2, TNF-receptor-associated factor 2; CHOP, C/EBP homologous protein.
ALT level was observed in the I/R group compared with that in the Sham group after 6 h of reperfusion and the ALT level returned to normal in the I/R group compared with that in the Sham group after 12 h reperfusion (Fig. 1). Pretreatment with NaHS prevented the elevations in ALT levels 6 h after I/R compared with those in the I/R group. A significant increase in the ALT level was observed in the I/R-PAG group after 6 and 12 h of reperfusion compared with those in the Sham group. These results indicated that endogenous H$_2$S may have a protective function in liver I/R, and exogenous H$_2$S ameliorates hepatic I/R injury.

H$_2$S ameliorates cell apoptosis in I/R and inhibition of endogenous H$_2$S results in aggravation of cell apoptosis. To investigate the functions of H$_2$S during liver I/R, the I/R-rats were pretreated with NaHS or the endogenous H$_2$S inhibitor, PAG. TUNEL assay was performed to assess the percentage of apoptotic cells in the livers of the rats of the four groups. As shown in Fig. 2, the hepatic cell apoptosis induced by I/R was decreased by NaHS intervention; however, the apoptosis rate was increased in the I/R-PAG group compared with the I/R group after 6 and 12 h of reperfusion. Next, total protein and mRNA were extracted from the livers and the expression levels of caspase-12, which is an apoptosis-promoting protease, were detected in the four groups. As shown in Figs. 3 and 4, caspase-12 was highly expressed in the I/R and I/R-PAG groups, while no difference in caspase-12 expression was found between the I/R-PAG and I/R groups. More importantly, NaHS intervention significantly alleviated I/R-induced caspase-12 upregulation.

H$_2$S inhibits ERS and inhibition of endogenous H$_2$S results in aggravation of ERS induced by I/R injury. To investigate the molecular mechanism underlying the protective effects of H$_2$S, the protein and mRNA levels of ERS-associated molecules were analyzed in liver tissue by western blot and...
RT-qPCR assays. As shown in Figs. 3 and 4, the protein and mRNA levels of ATF6, PERK, GRP78, TRAF2 and CHOP in the liver were markedly elevated by I/R, whereas NaHS treatment attenuated the stimulatory effect of I/R on ATF6, PERK, GRP78, TRAF2 and CHOP protein and mRNA levels. PAG treatment exerted no obvious effect on the action of I/R on the protein and mRNA levels of ATF6, PERK, GRP78, TRAF2 and CHOP compared with the I/R group; therefore, it was inferred that endogenous H$_2$S may be involved in pathways other than ERS-related pathways.

Figure 4. Reverse transcription-qPCR analysis was used to examine the mRNA levels of endoplasmic reticulum stress-associated genes. Each bar represents the mean ± SEM of three separate repeated experiments.  *P<0.05 vs. sham group;  #P<0.05 vs. 6 h;  $P<0.05 vs. I/R group. I/R, ischemia/reperfusion; PERK, pancreatic ER eukaryotic translation initiation factor-2α kinase; ATF6, activating transcription factor-6; GRP78, glucose-regulated protein 78; TRAF2, TNF-receptor-associated factor 2; CHOP, C/EBP homologous protein.
Discussion

The present study demonstrated that exogenous H$_2$S treatment, using NaHS as the H$_2$S donor, reduced I/R injury, whereas inhibition of endogenous H$_2$S by PAG treatment aggravated I/R injury in a hepatic I/R rat model. In accordance with histological results, serum ALT measurement demonstrated that the levels of ALT were elevated in the I/R and I/R-PAG groups, indicating that H$_2$S has a hepatoprotective function against I/R. Similar with previous findings, whereby Na2S was used as the H$_2$S donor, H$_2$S attenuated the elevation of ALT after I/R (14).

The present study further demonstrated that the protective effect of H$_2$S was associated with its attenuation of ERS induced by hepatic I/R injury, as the protein and mRNA expression of ERS-associated molecules, including ATF6, PERK, GRP78, TRAF2 and CHOP, were inhibited by NaHS treatment. A previous study also demonstrated that H$_2$S exerts protective effects on the cardiovascular system, as it attenuates cardiomyocytic ERS induced by hyperhomocysteinemia in rats (18). ERS-induced apoptosis may play a significant role in the pathophysiology of I/R injury (19,20).

ERS triggers the UPR pathway, which has three branches, involving IRE1, PERK and ARF6 (21). GRP78, an indicator of UPR pathway activation also referred to as immunoglobulin heavy chain-binding protein, is a central regulator of ER function due to its roles in protein folding and assembly, targeting of misfolded proteins for degradation, ER Ca$^{2+}$ binding and controlling the activation of transmembrane ERS sensors (21). Induction of GRP78 has been widely used as a marker for ERS and the onset of the UPR. Due to its anti-apoptotic properties, stress induction of GRP78 represents an important prosurvival component of the evolutionarily conserved UPR (22). Active IRE1 recruits the adaptor TRAF2, which activates apoptosis via the c-Jun N-terminal kinase pathway (23,24). CHOP was originally identified in response to DNA damage, and the CHOP level is a sensitive indicator of ERS (23). Caspase-12 is an enzyme that mediates ER-specific apoptosis and cytotoxicity (25,26) and it was found to be highly expressed in the I/R and I/R-PAG groups, while no difference in caspase-12 expression was found between the I/R-PAG and I/R groups. Therefore, caspase-12 may not play an important part in the process of apoptosis that is caused by inhibition of endogenous H$_2$S. A previous study also demonstrated that traumatic-hemorrhagic shock or reperfusion induces early and persistent ERS of the liver protein and nitrosylative stress (6,27). The present study demonstrated that NaHS treatment attenuated the stimulatory action of I/R on ATF6, PERK, GRP78, TRAF2 and CHOP protein and mRNA levels, which was consistent with the findings of the aforementioned previous studies.

Endogenous H$_2$S is a gaseous mediator produced by CSE in numerous tissues, including the liver (28). The enzymatic activity of CSE was found to be inhibited by PAG (29). To determine the contribution of endogenous H$_2$S to liver protection against I/R, PAG was used to inhibit the production of endogenous H$_2$S. Low levels of H$_2$S in the body result in severe damage following I/R injury, with marked elevation of ALT levels. However, the expression levels of ERS- and cell apoptosis-related genes from I/R-PAG rats were similar with those in I/R rats. Therefore, endogenous H$_2$S may also be involved in pathways other than ERS-related pathways.

In summary, the present study demonstrated that exogenous H$_2$S can effectively ameliorate I/R-induced injury along with decreased protein and mRNA expression of ERS-associated molecules, including ATF6, PERK, GRP78, TRAF2 and CHOP. I/R was shown to induce ERS and activate ERS-related apoptosis pathways, whereas inhibition of endogenous H$_2$S led to aggravation of damage after I/R and exogenous H$_2$S markedly reduced I/R injury. However, these conclusions were drawn based on the results of the hepatic I/R rat model constructed in the present study and they must be interpreted with caution, as the underlying molecular mechanisms require further investigation.

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Availability of data and materials

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

Authors’ contributions

HNF and XLW conceived the study. LC and KQM participated in all the experiments; XLW and TSC analyzed the data; LC and KQM have seen and approved the final version of this manuscript. All authors agree to be accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved.

Ethics approval and consent to participate

Animal studies were performed in compliance with the guidelines of animal ethics of Qinghai University and were approved by the Animal Care Committee of Qinghai University (no. 20180152).

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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