

Pyroptosis, a novel mechanism implicated in cataracts

XIN JIN¹, HAO JIN², YAN SHI¹, YIYUAN GUO¹ and HONG ZHANG¹

Departments of ¹Ophthalmology and ²Orthopedics, The First Affiliated Hospital of Harbin Medical University, Harbin, Heilongjiang 150001, P.R. China

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Abstract. An understanding of the mechanism of cataract formation may reduce its burden on medical care worldwide. It is established that pyroptosis is associated with oxidative stress, one of the causes of cataracts, and may provide novel therapeutic targets for the treatment of cataracts. The present study therefore investigated the role of pyroptosis in cataract formation. SRA01/04 human lens epithelium cells (HLECs) were treated with H₂O₂ and cell viability was assessed by an MTT assay. Pyroptosis in HLECs was examined by TUNEL staining, and the expression of caspase-1 and interleukin (IL)-1 β was determined using reverse transcription-quantitative polymerase chain reaction (RT-qPCR), western blot analysis and immunostaining. A caspase-1 inhibitor was used to investigate the effects of caspase-1 downregulation. In addition, the expression of caspase-1 and IL-1 β in lens anterior capsule tissue samples from patients with cataracts and normal controls was also analyzed by immunostaining, RT-qPCR and western blot analysis. The results demonstrated that pyroptosis in H₂O₂-treated HLECs, and the mRNA and protein expression of caspase-1 and IL-1 β , was significantly increased compared with control cells. Furthermore, caspase-1 and IL-1 β expression was significantly increased in cataract tissue samples compared with normal controls. When HLECs were cotreated with a caspase-1 inhibitor and 100 μ M H₂O₂, caspase-1 and IL-1 β expression were decreased compared with the 100 μ M H₂O₂-only group. In conclusion, the results of the present study demonstrate that pyroptosis may have a role in cataract formation, and the caspase-1 and IL-1 β pathways may be

involved in this pathological process. Pyroptosis appears to be a promising target in the prevention of cataract formation.

Introduction

As a global problem, cataracts are the leading cause of blindness in the elderly population and account for almost half of visually disabled people (1). Surgery has become an effective treatment method for cataracts. However, risks of surgical complications exist (2,3). Understanding the mechanisms of cataracts may allow the development of cataracts to be prevented, and subsequently reduce risks of surgical complications and the burden of cataracts on medical care worldwide.

In recent decades, emerging research has enriched our molecular understanding of the various forms of programmed cell death, which includes several endogenous genetically defined pathways (4). Apoptosis is the most widely recognized form of programmed cell death and involves particular caspases (cysteine-dependent aspartate-specific proteases), which lead to coordinated cell disassembly (5,6). It has been established that apoptosis of lens epithelial cells is a common cellular basis for the initiation and progression for non-congenital cataracts (7). Additional forms of programmed cell death include autophagy, oncosis and pyroptosis (also termed caspase 1-dependent programmed cell death) (8).

Pyroptosis is associated with rapid plasma membrane rupturing and the release of proinflammatory intracellular contents (9), and may be initiated by numerous pathological stimuli, including brain injury (10), myocardial infarction (11) or cancer (12), and has an important function in the control of microbial infections (13). Pyroptosis is closely associated with oxidative stress (14,15). Several studies have demonstrated that oxidative stress has a key role in cataractogenesis *in vivo* and *in vitro* (16,17). However, whether pyroptosis is implicated in the initiation and progression of non-congenital cataracts remains to be established.

The present study investigated the involvement of pyroptosis in cataract formation in a human lens epithelium cell line, and also characterized the expression of caspase-1 and interleukin (IL)-1 β in cataract lens anterior capsule tissue samples.

Materials and methods

Materials. The SV40 T-antigen-transformed human lens epithelial cell line (SRA01/04 cells) was obtained from the American Type Culture Collection (Manassas, VA, USA).

Correspondence to: Professor Hong Zhang, Department of Ophthalmology, The First Affiliated Hospital of Harbin Medical University, 23 Youzheng Road, Harbin, Heilongjiang 150001, P.R. China
E-mail: hydggzh@163.com

Abbreviations: HLECs, human lens epithelium cells; FBS, fetal bovine serum; DMEM, Dulbecco's modified Eagle's medium; RT-qPCR, reverse transcription-quantitative polymerase chain reaction; ROS, reactive oxygen species

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Fetal bovine serum (FBS) and Dulbecco's modified Eagle's medium/F12 (DMEM/F12) were obtained from Biological Industries (Kibbutz Beit-Haemek, Israel). H₂O₂ was purchased from Shanghai Zhongshi Chemistry Industry Co., Ltd. (Shanghai, China). MTT was purchased from Sigma-Aldrich (Merck KGaA, Darmstadt, Germany). TRIzol and polymerase chain reaction (PCR) primers were purchased from Invitrogen (Thermo Fisher Scientific, Inc., Waltham, MA, USA). The NanoDrop spectrophotometer was obtained from NanoDrop Technologies (Thermo Fisher Scientific, Inc.). Primary antibodies were purchased from Santa Cruz Biotechnology, Inc. (Dallas, TX, USA). Fluorochrome-labeled secondary antibody (Alexa Fluor 800) was obtained from Thermo Fisher Scientific, Inc.

Human tissue samples. A total of 40 cataract lens anterior capsule samples were obtained from patients with cataracts that were free from other types of ocular diseases (age range, 68-80 years) in the operation room of the Eye Hospital of Harbin Medical University (Harbin, China), between October 2015 and March 2016. In addition, 40 control lens anterior capsular samples were collected from healthy donor eyes (age range, 54-69 years) that were free of any ocular diseases and donated to the Eye Bank of Heilongjiang Province (Harbin, China), between July 2014 to March 2016. Samples were either embedded in paraffin or immediately snap-frozen and stored at -80°C until RNA extraction. All samples were collected with informed consent and the study was approved by the Research Ethics Committee of Harbin Medical University. Patient information is listed in Table I.

Cell culture. SRA01/04 cells (1x10⁵) were cultured in DMEM/F12 supplemented with 20% FBS at 37°C in 5% CO₂ overnight. Cells in the logarithmic growth phase were collected and treated with 0, 25, 50 or 100 μM H₂O₂ for 0, 24 or 48 h at 37°C in 5% CO₂. Control cells were treated with DMEM/F12 containing 20% FBS alone. Caspase-1 inhibitor for down-regulation was synthesized by Cayman Chemical Company (item no. 10014; Ann Arbor, MI, USA). After the cell density reached 80%, 100 μM caspase-1 inhibitor was added simultaneously with H₂O₂ to the designated well. After 0, 24 or 48 h of treatment at 37°C, cells were harvested for subsequent experiments.

Cell proliferation assay. MTT were used in accordance with the manufacturer's protocol. SRA01/04 cells were seeded in 96-well plates at 1x10⁴ cells/well and treated with 0, 25, 50, 100 or 200 μM H₂O₂ and maintained for 24 or 48 h at 37°C. MTT solution (10 μl) was added to each well and cells were incubated in 37°C for 2 h. DMSO (150 μl) was added to each well. The absorbance at a wavelength of 450 nm was evaluated using a microplate reader. The data are representative of three individual experiments in triplicate.

Reverse transcription-quantitative PCR (RT-qPCR). Caspase-1 and IL-1β mRNA expression was determined by RT-qPCR using SYBR Green detection reagents. Primer sequences are listed in Table II. Total RNA was extracted from each sample with TRIzol reagent, according to the manufacturer's protocol. Total isolated RNA (500 ng) was used to synthesize cDNA

in a 20 μl reaction mixture containing 2 μl RNA, 4 μl 5X RT buffer, 1 μl primer mix, 1 μl RT enzyme and 12 μl H₂O (Toyobo Life Science, Osaka, Japan). qPCR amplification was performed in a 20 μl reaction volume containing 2 μl cDNA, 6 μl diethyl pyrocarbonate, 10 μl SYBR Master mix (Toyobo Life Science), 1 μl forward primer and 1 μl reverse primer. qPCR was performed using the ABI PRISM 7500 Sequence Detection System (Applied Biosystems; Thermo Fisher Scientific, Inc.) and PCR conditions were as follows: 95°C for 60 sec, followed by 40 cycles of 95°C for 15 sec, 60°C for 15 sec and 72°C for 45 sec. The housekeeping gene GAPDH was used as an internal positive control standard for quantitative analysis in triplicate. Cq values obtained via the 2^{-ΔΔCq} method (18) were used to quantify mRNA expression.

Western blot analysis. Western blot analysis was performed to detect the expression levels of certain proteins of interest. Samples were lysed using radioimmunoprecipitation assay buffer (Beijing Solarbio Science & Technology Co., Ltd., Beijing, China). A bicinchoninic acid assay (Beijing Solarbio Science & Technology Co., Ltd.) was used to determine the protein concentration of the samples. Protein (50 μg) per sample was separated using 10% SDS-PAGE and transferred onto nitrocellulose membranes. Membranes were blocked in 5% milk for 2 h at room temperature, and then immunoblotted overnight with rabbit caspase-1 (1:1,000; cat. no. PRS3459; Sigma-Aldrich; Merck KGaA), IL-1β (1:1,000; cat. no. SAB4503272; Sigma-Aldrich; Merck KGaA) and GAPDH (1:800; cat. no. ab8245; Abcam, Cambridge, UK) primary antibodies at 4°C with gentle shaking. Subsequently, membranes were incubated with goat anti-rabbit fluorochrome-labeled secondary antibody for 2 h at room temperature (1:5,000; Alexa Fluor 800; cat. no. A32735). Immunoreactivity was detected with an Odyssey fluorescent scanning system (LI-COR Biosciences, Lincoln, NE, USA) and analyzed by Image Studio software version 4.0 (LI-COR Biosciences). GAPDH was used as loading control.

Immunofluorescence and immunohistochemistry staining. For immunofluorescence staining, SRA01/04 cells were seeded in 24-well plates at a density of 3x10⁵ cells/well and maintained for 24 h. Following treatment with 100 μM H₂O₂ with or without caspase-1 inhibitor treatment, cells were fixed with 4% paraformaldehyde for 1 h at 37°C, then permeabilized with 0.1% Triton for 15 min at room temperature. After rinsing with PBS, cells were blocked with 10% bovine serum albumin (Biosharp, China) 30 min at room temperature. The cells were subsequently incubated with rabbit caspase-1 (1:200) or IL-1β (1:200) antibodies overnight at 37°C. Cells were rinsed three times in PBS and incubated with Cy3-conjugated goat anti-rabbit IgG (1:100; cat. no. AP132C; Sigma-Aldrich; Merck KGaA) secondary antibody 30 min at room temperature. Staining was observed using an Observer A1 fluorescence microscope (Zeiss GmbH, Jena, Germany), and ZEN software version 2 (Zeiss GmbH) was used.

For immunohistochemistry staining, lens anterior capsule tissue samples were placed on cover slides and fixed with 4% paraformaldehyde for 30 min at room temperature. After rinsing with PBS, capsular samples were penetrated with 0.1% Triton X-100 15 min and blocked with 5% BSA 30 min at room

Table I. Patient and volunteer information.

Tissue sample type	Mean age, years (\pm SD)	Age range, years	Males (%)	Females (%)
Cataracts, n=40	74.4 (\pm 3.84)	68-80	17 (42.5)	23 (57.5)
Normal, n=40	62.4 (\pm 4.04)	54-69	20 (50)	20 (50)

SD, standard deviation.

Table II. Primers used for reverse transcription-quantitative polymerase chain reaction.

Gene	Primer sequence	
	Forward	Reverse
GAPDH	5'-AAGAAGGTGGTGAAGCAGGC-3'	5'-TCCACCACCCTGTTGCTGTA-3'
Caspase-1	5'-ACACGTCTTGCCCTCATTATCT-3'	5'-ATAACCTGGGCTTGTCTTTCA-3'
Interleukin-1 β	5'-CCTTGTCGAGAATGGGCAGT-3'	5'-TTCTGTCGACAATGCTGCCT-3'

temperature. Slides were subsequently incubated with primary antibodies against caspase-1 (1:200) and IL-1 β (1:200) at 4°C overnight. Slides were then washed using PBS and further incubated with secondary antibodies conjugated to Cy3 (1:100; cat no. AP132C; Sigma-Aldrich; Merck KGaA) for 2 h at room temperature, followed by staining with 3,3'-diaminobenzidine 15 sec at room temperature. Samples were hydrated in graded ethanol. Staining was observed using an Observer A1 fluorescence microscope (Zeiss GmbH, Jena, Germany), and ZEN software version 2 (Zeiss GmbH) was used.

TUNEL staining. SRA01/04 cells (3×10^5 cells/well) were plated onto coverslips in 24-well culture plates and an *in situ* Cell Death Detection kit (Fluorescein; Roche, Diagnostics; Indianapolis, IN, USA) was employed to detect DNA fragmentation of individual cells, according to the manufacturer's protocol. The nuclei were stained with DAPI (1 μ g/ml) for 5 min at room temperature. TUNEL staining was assessed by fluorescence microscopy (Eclipse 80i; Nikon Co., Tokyo, Japan) at x200 magnification in 6 fields of view. Nuclei that were double labeled with DAPI and TUNEL were considered positive.

Statistical analysis. Statistical significance was determined by a two-tailed Student's t-test or one-way analysis of variance followed by Tukey's post hoc test. Data are presented as the mean \pm standard deviation. The results were analyzed by SPSS 18.0 software (SPSS, Inc., Chicago, IL, USA) and $P < 0.05$ was considered to indicate a statistically significant difference.

Results

Pyroptosis is induced in H₂O₂-treated SRA01/04 cells. Caspase-1 and IL-1 β are established markers of pyroptosis. To determine whether pyroptosis is involved in cataract formation, SRA01/04 lens epithelial cells were exposed to various concentrations of H₂O₂ (0, 25, 50 and 100 μ M) for 24 or 48 h. Pyroptosis was detected by morphology analyses, MTT, RT-qPCR and western blot assays, and TUNEL staining

(Figs. 1 and 2, respectively). SRA01/04 cells became swollen and the cell number decreased with increasing H₂O₂ concentration. MTT assay results demonstrate that SRA01/04 cell growth was inhibited by 25, 50, 100 and 200 μ M H₂O₂ at 24 and 48 h (Fig. 1B). The mRNA and protein expression of caspase-1 and IL-1 β were increased in H₂O₂-treated SRA01/04 cells in a dose-dependent manner (Fig. 1C-F). Compared with the control group, relative caspase-1 and IL-1 β expression increased >50% when the H₂O₂ concentration was 100 μ M. Furthermore, TUNEL assay results provided further evidence that H₂O₂-induced cell pyroptosis was significantly increased in a dose-dependent manner (Fig. 2). Based on these results, 100 μ M H₂O₂ was selected for subsequent experiments.

Validation of pyroptosis in the lens anterior capsules of patients with cataracts. To further validate these results, cataract lens anterior capsules and normal lens anterior capsule samples were collected and analyzed for caspase-1 and IL-1 β expression using immunohistochemistry staining, RT-qPCR and western blot analysis. Immunohistochemistry staining results demonstrated increased positive staining of caspase-1 and IL-1 β in the cataract lens anterior capsular specimens compared with normal samples (Fig. 3A). The mRNA and protein expression levels of caspase-1 and IL-1 β were also significantly upregulated in cataract samples compared with normal anterior lens capsules (Fig. 3B-E).

Caspase-1 inhibitor reduces pyroptosis in SRA01/04 cells. To confirm the role of caspase-1 in pyroptosis in SRA01/04 cells, caspase-1 was downregulated by an irreversible inhibitor. Caspase-1 inhibitor was incubated with SRA01/04 cells and the efficacy was confirmed by immunofluorescence staining, RT-qPCR and western-blot analysis (Fig. 4A-C). In addition, pyroptosis was significantly reduced by caspase-1 inhibitor treatment compared with the 100 μ M H₂O₂-only group, as demonstrated by TUNEL staining (Fig. 4D). Furthermore, the mRNA and protein expression of IL-1 β , which functions downstream of caspase-1, was significantly decreased in

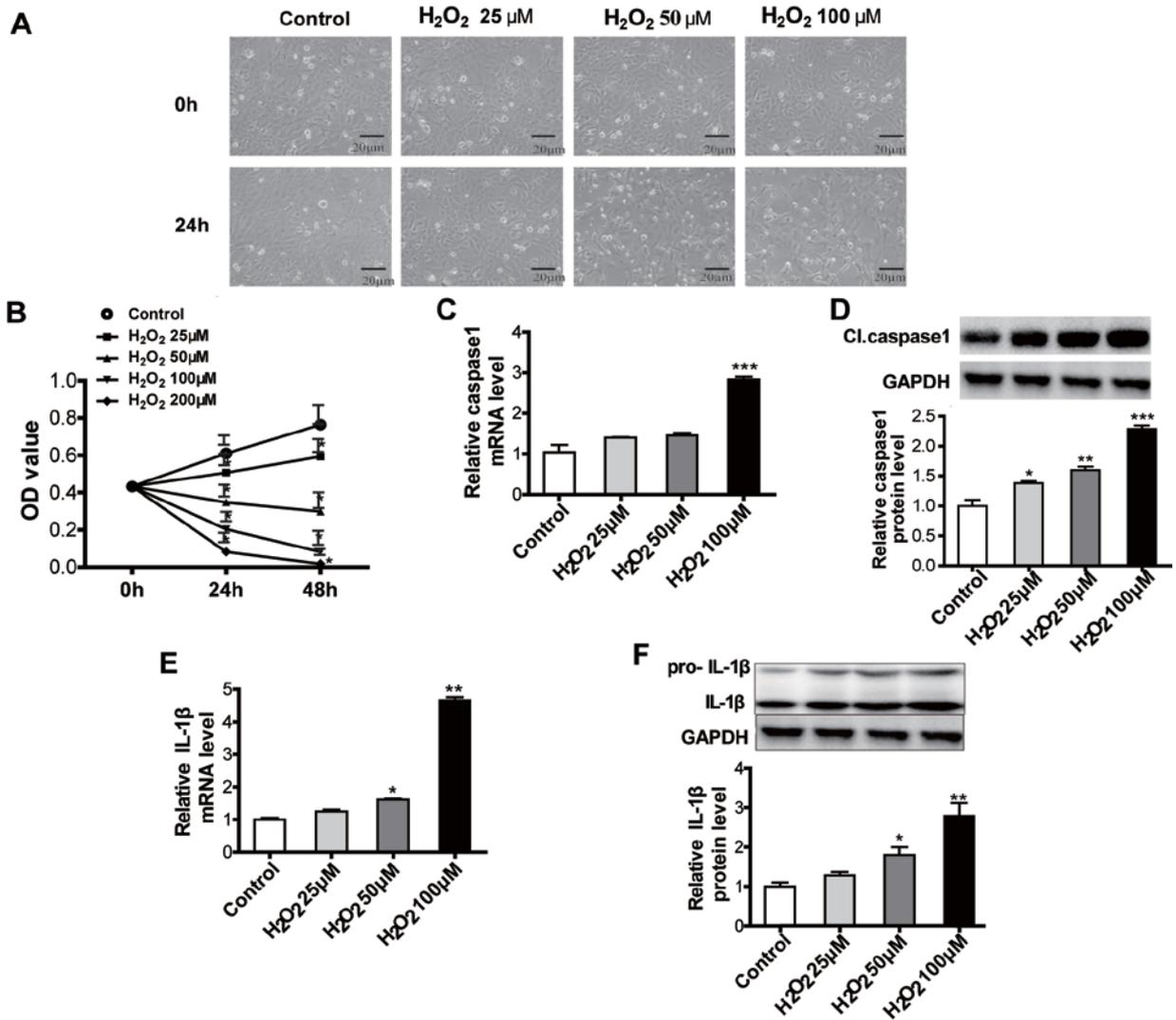


Figure 1. Pyroptosis was increased in H₂O₂ treated human lens epithelium cells. (A) When exposed to 0, 25, 50 and 100 μM H₂O₂ for 24 h, SRA01/04 cells became swollen and the cell number decreased with increasing H₂O₂ concentration. Scale bar, 20 μm. x200 magnification. (B) MTT assay results demonstrated that SRA01/04 cell growth was inhibited when treated with 25, 50, 100 and 200 μM H₂O₂ for 24 and 48 h. (C) RT-qPCR results indicated that relative caspase-1 mRNA levels were increased with increasing concentrations of H₂O₂. (D) Cl. caspase-1 protein levels were upregulated increasing H₂O₂ concentrations, as demonstrated by western blot analysis. (E) RT-qPCR results demonstrated that relative IL-1β mRNA levels were increased with increasing concentrations of H₂O₂. (F) IL-1β protein levels were upregulated with increasing H₂O₂ concentrations, as demonstrated by western blot analysis. *P<0.05, **P<0.01 and ***P<0.001 vs. control. RT-qPCR, reverse transcription-quantitative polymerase chain reaction; Cl., cleaved; IL, interleukin; OD, optical density; pro-, precursor.

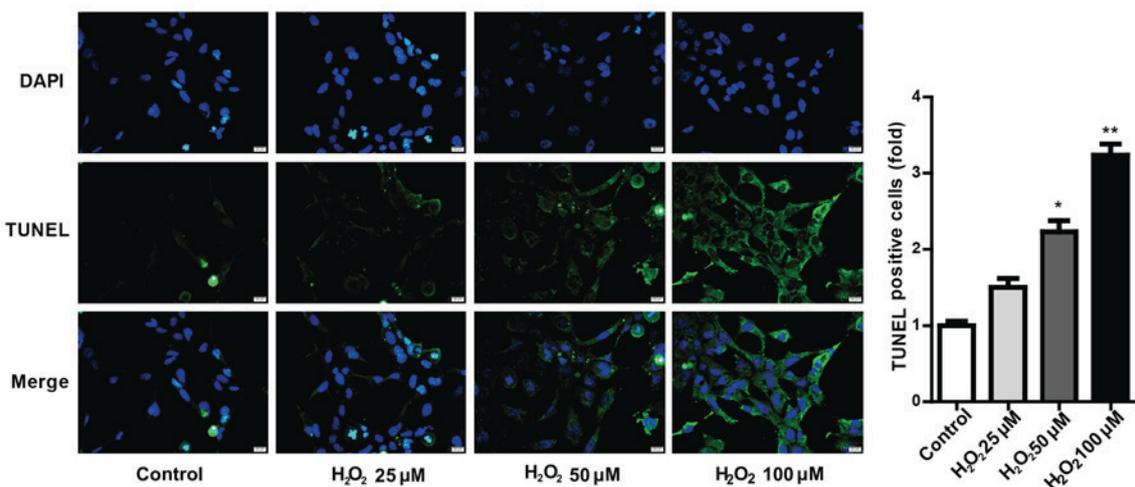


Figure 2. TUNEL staining images of SRA01/04 cells. Cells were treated with 0, 25, 50 and 100 μM H₂O₂ and TUNEL analysis was performed. Blue, nuclear staining with DAPI; green, TUNEL staining. Scale bar, 20 μm.

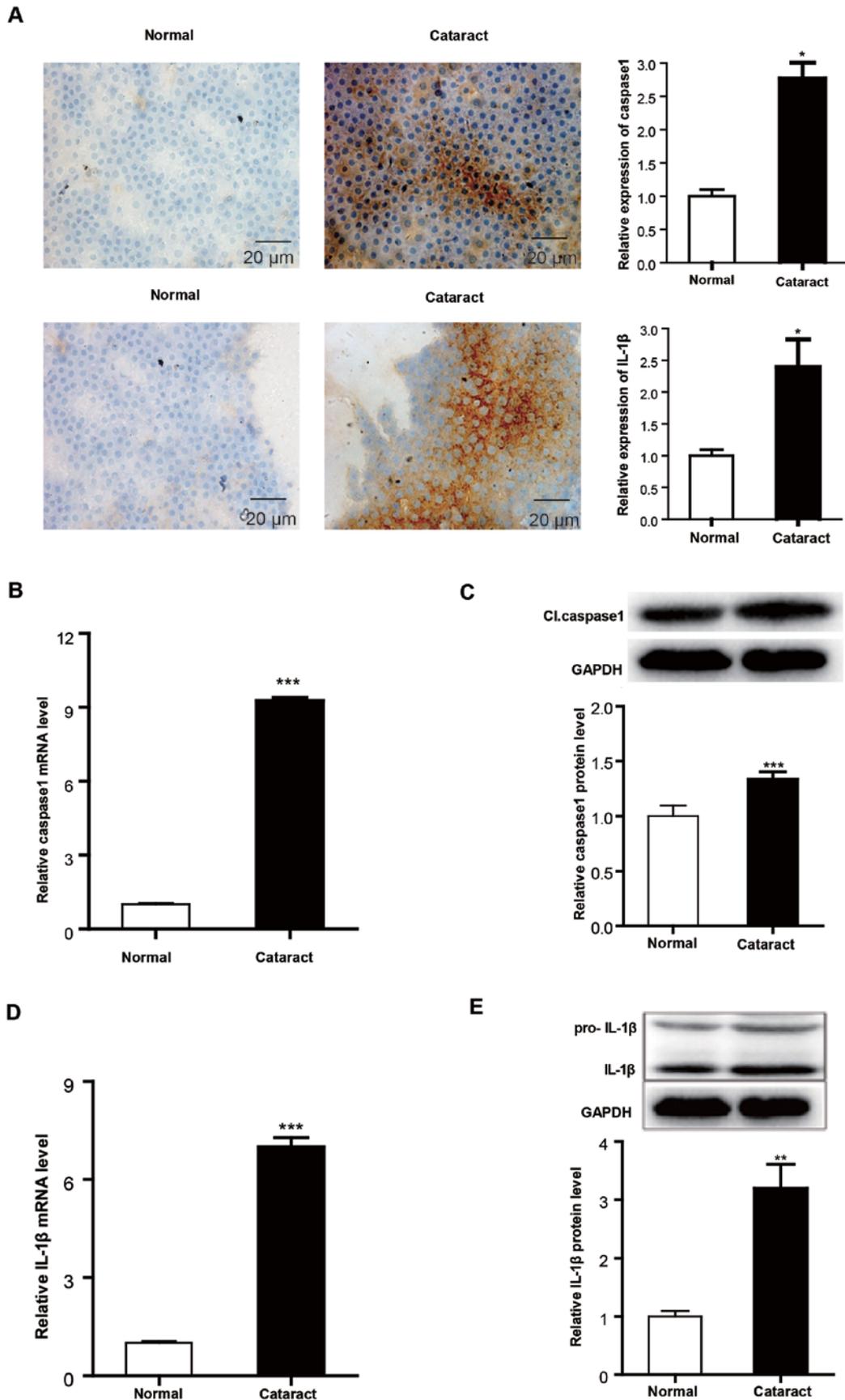


Figure 3. Validation of pyroptosis in the lens anterior capsules of patients with cataracts. (A) Immunohistochemical staining of caspase-1 and IL-1 β in representative anterior lens capsule tissue specimens compared with normal controls. Scale bar, 20 μ m. (B) RT-qPCR results of the relative mRNA levels of caspase-1 in anterior lens capsules of normal donors and patients with cataracts. (C) Western blot analysis of cl. caspase-1 protein levels in anterior lens capsules of normal donors and patients with cataracts. (D) RT-qPCR results of the relative mRNA levels of IL-1 β in anterior lens capsules of normal donors and patients with cataracts. (E) Western blot analysis of IL-1 β protein levels in anterior lens capsules of normal donors and patients with cataracts. * P <0.05, ** P <0.01 and *** P <0.001 vs. normal donors. IL, interleukin; RT-qPCR, reverse transcription-quantitative polymerase chain reaction; cl., cleaved; pro-, precursor.

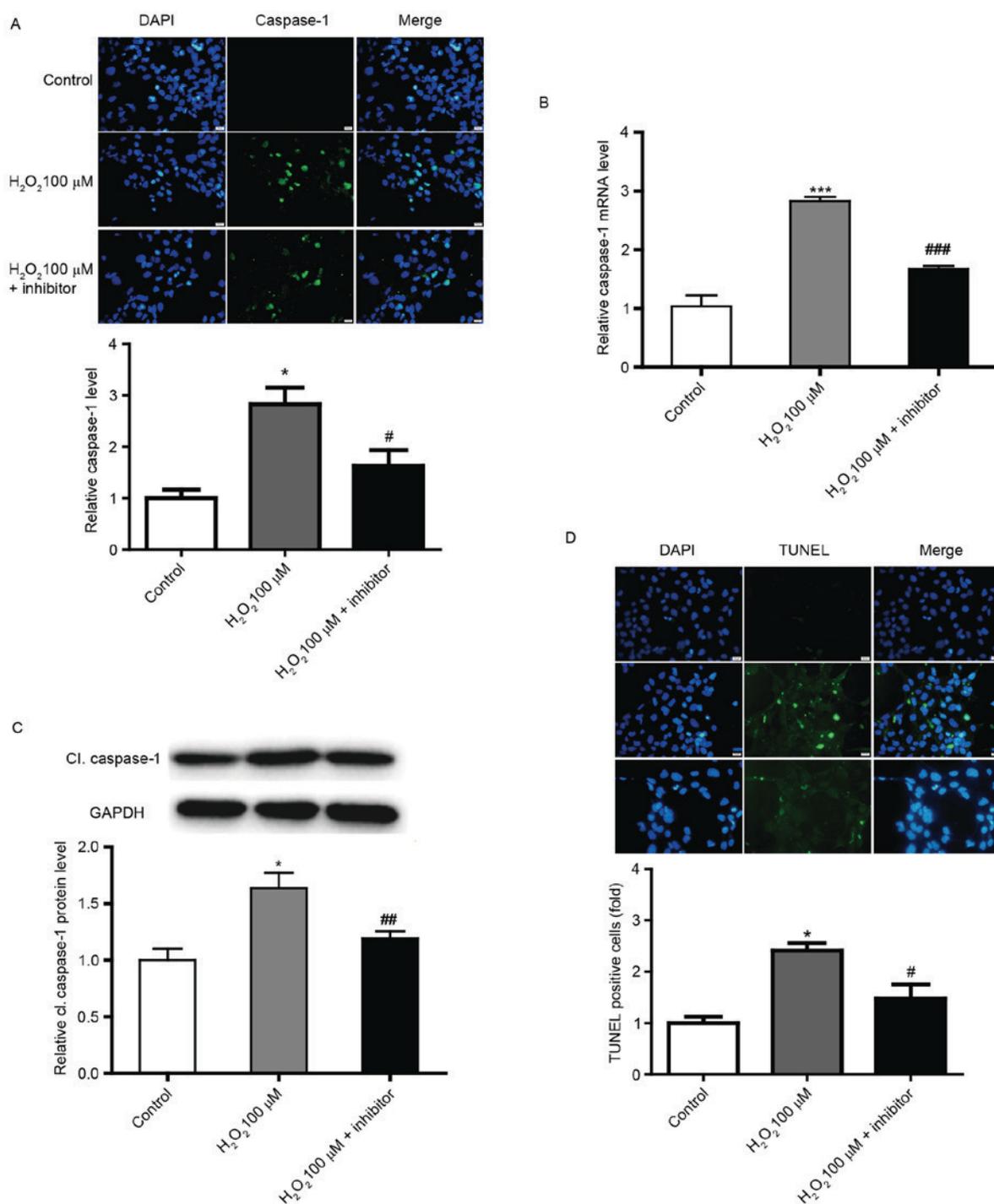


Figure 4. Downregulation of caspase-1 in human lens epithelium cells. (A) Immunofluorescence images demonstrating the expression of caspase-1 in SRA01/04 cells in control, 100 μ M H₂O₂ and 100 μ M H₂O₂ + caspase-1 inhibitor groups. Blue, nuclear staining with DAPI; green, caspase-1 staining. Scale bar, 20 μ m. Relative (B) mRNA levels of caspase-1 and (C) protein levels of cl. caspase-1 in SRA01/04 cells in control, 100 μ M H₂O₂ and 100 μ M H₂O₂ + caspase-1 inhibitor groups. (D) TUNEL staining images of SRA01/04 cells in control, 100 μ M H₂O₂ and 100 μ M H₂O₂ + caspase-1 inhibitor groups. Blue, nuclear staining with DAPI; green, TUNEL staining. Scale bar, 20 μ m. * P <0.05 and *** P <0.001 vs. control; # P <0.05, ## P <0.01 and ### P <0.001 vs. 100 μ M H₂O₂-only group. cl., cleaved.

caspase-1 inhibitor-treated SRA01/04 cells compared with the 100 μ M H₂O₂ only group (Fig. 5).

Discussion

In recent years, increasing studies have focused on understanding the mechanisms of pyroptosis in various diseases,

and identifying the genes and pathways that are implicated in this process. The present study reported that pyroptosis may have an important role in human lens epithelial cells under oxidative stress. To investigate the effect of pyroptosis in cataract formation, the present study demonstrated that caspase-1 expression was increased in lens epithelium cells treated with H₂O₂. In addition, the results also confirmed that caspase-1

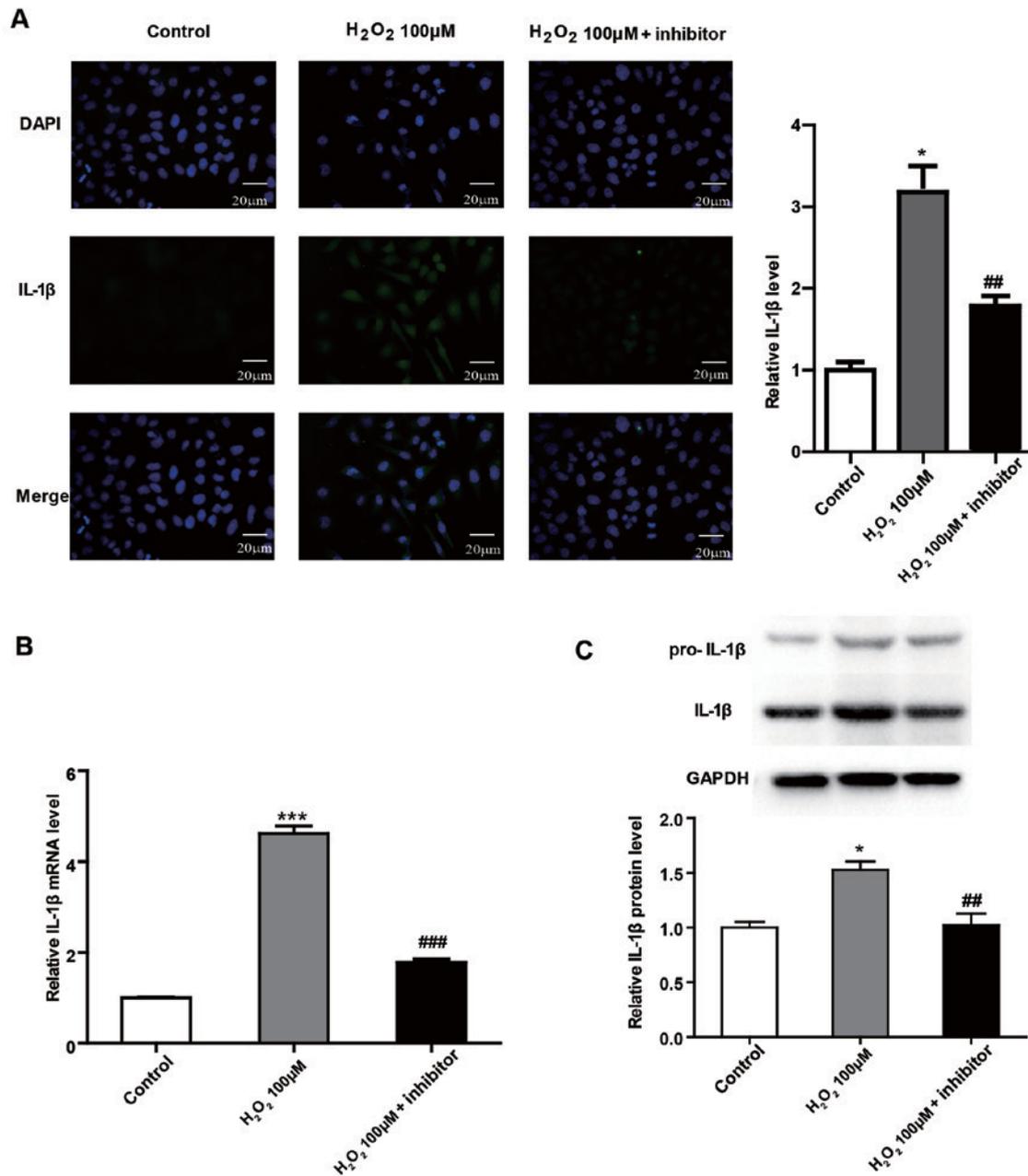


Figure 5. Caspase-1 inhibitor treatment reduced pyroptosis in human lens epithelium cells. (A) Immunofluorescence images demonstrating the expression of IL-1β in SRA01/04 cells in control, 100 μM H₂O₂ and 100 μM H₂O₂ + caspase-1 inhibitor groups. Blue, nuclear staining with DAPI; green, IL-1β staining. Scale bar, 20 μm. Relative (B) mRNA and (C) protein levels of IL-1β in SRA01/04 cells in control, 100 μM H₂O₂ and 100 μM H₂O₂ + caspase-1 inhibitor groups. *P<0.05 and ***P<0.001 vs. control; **P<0.01 and ###P<0.001 vs. 100 μM H₂O₂-only group. IL, interleukin; pro-, precursor.

was significantly upregulated in cataract lens anterior capsule samples compared with normal lens anterior capsule samples. Furthermore, pyroptosis was decreased when caspase-1 was downregulated by using a caspase-1 inhibitor. Due to the relative low knockout efficiency of caspase-1 by the inhibitor that was designed and used in the present study, certain limitations are associated with our conclusions and further experimental verification is required.

Lens epithelial cells have a key role in stabilizing the intracellular environment and maintaining a clear crystalline lens. Consistent with various other degenerative ocular diseases, cataract formation occurs when the production rate of reactive oxygen species (ROS) exceeds the removal rate (19,20).

When lens epithelial cells are exposed to endogenous and exogenous oxidative stress, including growth factors, UVB radiation and inflammatory cytokines, they suffer oxidative injury and produce large amounts of ROS. Activation of ROS generation has been reported to induce NLR family pyrin domain containing 3/caspase-1 activation, which subsequently triggers IL-1β/IL-18 production and cell death by pyroptosis and apoptosis, in astroglial cells (21). H₂O₂ contains activated oxygen, which permeates cellular membranes and causes injury inside cells. H₂O₂ is commonly used for *in vitro* cellular oxidative damage models, such as cataract formation (22,23). The present study employed H₂O₂ to investigate the effects of pyroptosis in cataracts.

Caspase-1 and IL-1 β are important markers in the process of pyroptosis (24). The caspase family constitutes a family of cysteinyl aspartate proteases (25). Caspase-1 was the first caspase to be identified in mammalian cells and has an important function in apoptosis. Caspase-1 also mediates proinflammatory programmed cell death, also termed pyroptosis, in response to exogenous and endogenous stimuli to protect cells. Caspase-1 is not only activated in immune cells, but is also activated in epithelial and mesenchymal cells (26), and dysfunction of caspase-1 is closely associated with various diseases (27,28). However, whether pyroptosis of lens epithelial cells is implicated in the initiation and progression of non-congenital cataracts in humans remains to be established. The results of the current study demonstrate that caspase-1 expression was increased in H₂O₂-treated lens epithelial cells in a dose-dependent manner, as determined by RT-qPCR and western blot analysis. These results were also validated in the lens anterior capsules of cataract patients compared with normal lens anterior capsule samples. Although the increased levels of caspase-1 may indicate the involvement of pyroptosis in the process, it is difficult to distinguish between apoptosis and pyroptosis as apoptosis has also been reported to be involved in cataract formation in H₂O₂-treated lens epithelial cells (29). Therefore, further studies are required to address this issue.

Caspase-1 was initially characterized as a protease that converts the inactive precursors of IL-1 β and IL-18 (pro-IL-1 β /IL-18) into mature inflammatory cytokines, and was originally termed interleukin IL-1 β -converting enzyme (30). The present study also observed IL-1 β to be downstream of caspase-1 in SRA01/04 cells. When caspase-1 inhibitor was introduced to 100 μ M H₂O₂-treated SRA01/04 cells, IL-1 β expression was markedly decreased. The superfamily of IL-1 (IL-18, IL-1 β , pro-IL-18 and pro-IL-1 β) are the main proinflammatory cytokines inside the cell. Activated caspase-1 leads to the activation of pro-IL-1 β , and IL-1 β subsequently activates downstream nuclear factor- κ B (NF- κ B) signaling, which enhances the release of inflammatory cytokines (31). It has been reported that NF- κ B signaling is implicated in cataract formation (32). We hypothesize that H₂O₂-induced oxidative stress may activate NF- κ B signaling in human lens epithelial cells via caspase-1 activation and maturation of IL-1 β , which subsequently contributes to the development of cataracts. Further investigation is required to confirm this hypothesis. The authors of the present study plan to explore the association between NF- κ B signaling and pyroptosis by IL-33/sT2 signalling. IL-33 stimulation may include ERK, p38 MAPK and JNK as well as the activation of NF- κ B. It has been reported that caspase-1 can result in the release of IL-33 (33). It may be possible for future studies to explore this phenomenon in cataracts.

In conclusion, the results of the current study provide evidence that pyroptosis participates in the oxidation of human lens epithelial cells and may be involved in the initiation and progression of non-congenital cataracts. The results indicate a potential role of pyroptosis in cataract formation, which enhances our understanding of cataracts and may provide novel effective therapeutic methods for this condition.

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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

XJ analyzed and interpreted the patient data regarding the cataracts and was a major contributor in writing the manuscript. HJ and YS performed the histological examination of the lens anterior capsule samples. HZ and YG were involved in the design of the study. All authors read and approved the final manuscript.

Ethics approval and consent to participate

All samples were collected with informed consent and the study was approved by the Research Ethics Committee of Harbin Medical University.

Consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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