



# Hsa\_circ\_0003204 Knockdown Weakens Ox-LDL-Induced Cell Injury by Regulating miR-188-3p/TRPC6 Axis in Human Carotid Artery Endothelial Cells and THP-1 Cells

## OPEN ACCESS

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**Background:** Circular RNAs (circRNAs) are involved in atherosclerosis (AS) development. However, the function and mechanism of circRNA hsa\_circ\_0003204 (circ\_0003204) in carotid artery AS remain unclear.

**Methods:** Oxidized low-density lipoprotein (ox-LDL)-treated human carotid artery endothelial cells (HCtAECs) and THP-1 cells were used as cell models of carotid artery AS. Relative levels of circ\_0003204, microRNA-188-3p (miR-188-3p), and transient receptor potential canonical channel 6 (TRPC6) were detected by quantitative reverse transcription-polymerase chain reaction or Western blotting. The targeting relationship between circ\_0003204 or TRPC6 and miR-188-3p was assessed via dual-luciferase reporter analysis and RNA immunoprecipitation. Cell proliferation was assessed via 3-(4,5-dimethyl-2-thiazolyl)-2,5-diphenyl-2H-tetrazolium bromide assay and 5-ethynyl-2'-deoxyuridine (EdU) assay. Cell apoptosis was analyzed via assessing cell caspase-3 activity, apoptosis, and apoptosis-related protein. Inflammatory response was analyzed via analysis of interleukin-1 $\beta$  (IL-1 $\beta$ ), IL-6, and tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ). Oxidative stress was assessed via determination of reactive oxygen species (ROS), malondialdehyde (MDA), and superoxide dismutase (SOD).

**Results:** Circ\_0003204 and TRPC6 levels were elevated, and miR-188-3p expression declined in ox-LDL-treated HCtAECs and THP-1 cells. Circ\_0003204 could regulate TRPC6 expression via mediating miR-188-3p. Circ\_0003204 silencing weakened ox-LDL-induced viability inhibition and apoptosis in HCtAECs, and inflammatory response and oxidative stress in THP-1 cells via regulating miR-188-3p. MiR-188-3p overexpression attenuated ox-LDL-induced injury in HCtAECs and THP-1 cells by targeting TRPC6.

**Conclusion:** Circ\_0003204 knockdown mitigated ox-LDL-induced injury in HCTaECs and THP-1 cells via regulating the miR-188-3p/TRPC6 axis, indicating that circ\_0003204 might play an important role in carotid artery AS.

**Keywords:** carotid artery AS, ox-LDL, circ\_0003204, MiR-188-3p, TRPC6

## INTRODUCTION

Atherosclerosis (AS) is an inflammatory-related cardiovascular disease (1). Carotid artery AS is a group of AS and associated with the increased risk of cardiovascular disorders (2). Oxidized low-density lipoprotein (ox-LDL) has an essential role in AS development via regulating the function of multiple cell lines, like endothelial cells and macrophages (3). The malfunction of endothelial and THP-1 cells is implicated in the pathobiology of AS (4, 5). Therefore, analyzing the pathogenesis of ox-LDL-triggered dysfunction of carotid artery endothelial cells and THP-1 cells may help to explore new strategies for carotid artery AS treatment.

Circular RNAs (circRNAs) and microRNAs (miRNAs) are correlated with the regulation of cardiovascular cell biology in AS (6). CircRNAs are stable ncRNAs formed via back-splicing events, which act as vital biomarkers for cardiovascular diseases, including AS (7). CircRNA hsa\_circ\_0003204 (circ\_0003204), derived from ubiquitin-specific peptidase 36 (USP36), has been reported to be upregulated in ox-LDL-challenged human aortic endothelial cells (HAECs) and plays a vital role in cerebrovascular atherogenesis progression (8). Moreover, circ\_0003204 is reported to be dysregulated in ox-LDL-irritated human umbilical vein endothelial cells (HUVECs) (9). Hence, we assumed that circ\_0003204 might play a vital role in AS progression. However, how and whether circ\_0003204 takes part in the development of carotid artery AS remain unknown.

MiRNAs have been suggested to participate in AS development (10). According to the ceRNA hypothesis, circRNAs can regulate gene expression via binding to miRNAs (11). In this study, we screened the top 10 miRNAs (miR-1224-3p, miR-1236, miR-346, miR-370, miR-432, miR-593, miR-635, miR-1827, miR-620, and miR-188-3p) based on context + score percentage after circular RNA Interactome prediction. And preliminary experiments exhibited that miR-188-3p was highly pulled down by circ\_0003204 probe. A previous study indicated that miR-188-3p could inhibit the inflammatory response in AS mice (12). However, whether miR-188-3p is involved in circ\_0003204-mediated carotid artery AS development is unclear.

The transient receptor potential canonical channels (TRPCs) play an important role in cardiovascular diseases (13). Among all miR-188-3p targets predicted by the DIANA tool, five mRNAs [KLF6, IGF2, AKT3, TRIM14, and TRPC 6 (TRPC6)] have been reported to have the opposite function of miR-188-3p. And preliminary experiments showed that TRPC6 was pulled down the most by the miR-188-3p probe. TRPC6, a key member of TRPCs, contributes to ox-LDL-induced HAEC apoptosis (14). At present, the involvement of circ\_0003204, miR-188-3p, and

TRPC6 in carotid artery AS is unclear. Hence, we hypothesized that circ\_0003204 might regulate carotid artery AS development via mediating miR-188-3p/TRPC6 axis.

In this research, we utilized ox-LDL-challenged HCTaECs and THP-1 cells to mimic carotid artery AS environment (15). Moreover, we detected the expression of circ\_0003204, miR-188-3p, and TRPC6, and we explored the function of circ\_0003204 in ox-LDL-triggered cell injury. Additionally, we explored the ceRNA network of circ\_0003204/miR-188-3p/TRPC6 axis.

## MATERIALS AND METHODS

### Cell Culture and Treatment

HCTaECs were provided via Cell Applications (San Diego, CA, USA) and cultured in specific MesoEndo Cell Grown Medium (Cell Applications) in 5% CO<sub>2</sub> at 37°C. THP-1 cells were offered by Procell (Wuhan, China) and maintained in RPMI-1640 medium (Procell) containing 10% fetal bovine serum (FBS) (HyClone, Logan, UT, USA) and 1% penicillin/streptomycin (Thermo Fisher, Waltham, MA, USA) in 5% CO<sub>2</sub> at 37°C.

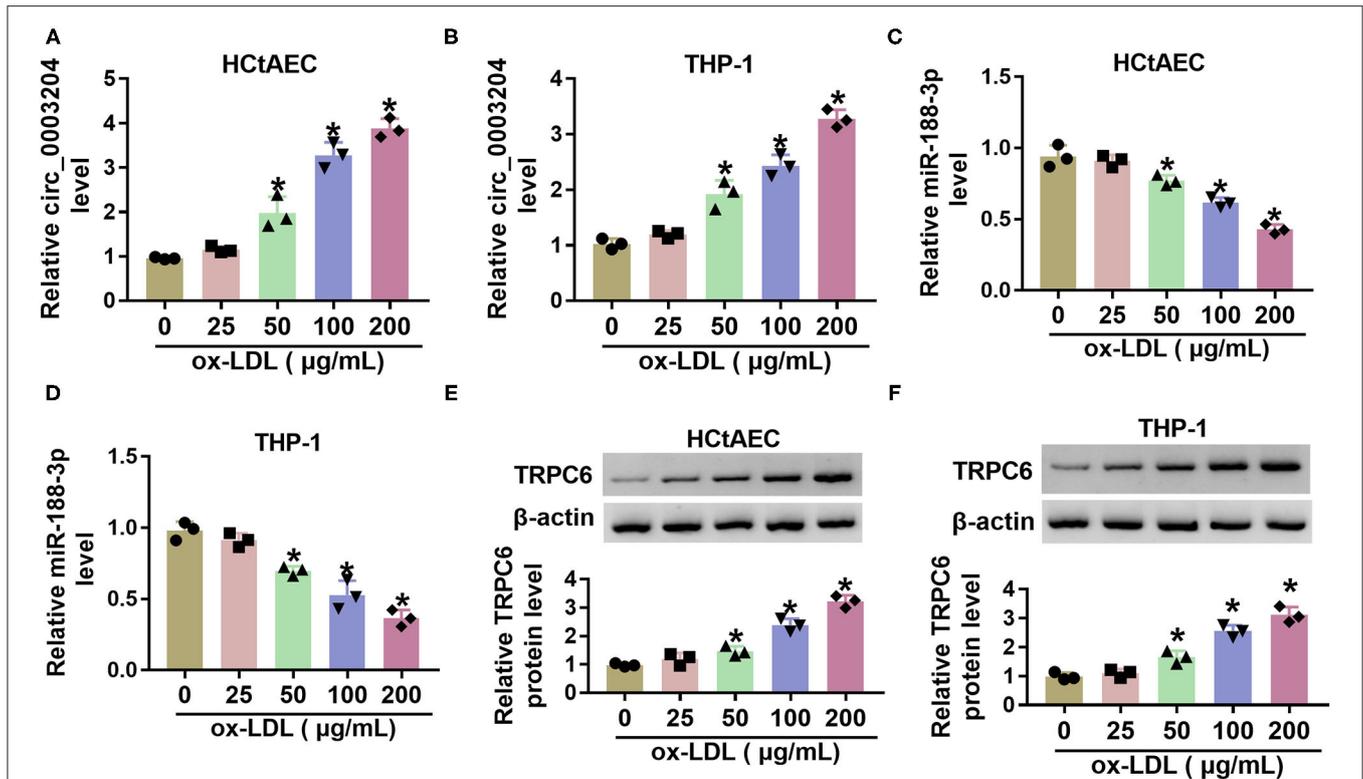
To mimic the carotid AS-like microenvironment, HCTaECs and THP-1 cells were challenged with different doses of ox-LDL (Solarbio, Beijing, China) for 24 h.

### Real-Time Quantitative PCR

The RNA was extracted by TRIzol (Thermo Fisher) (16). The RNA was reversely transcribed using a specific reverse transcription kit (Thermo Fisher). The cDNA was mixed with SYBR (Vazyme, Nanjing, China) and specific primers (Genscript, Nanjing, China) and used for qRT-PCR. The primers were as follows: circ\_0003204 (F, 5'-CTCAAATGCCCAAGGAGT GC-3'; R, 5'-GCAGGCGGCTGGATGATT-3'), TRPC6 (F, 5'-AGGGCTGGAGAGTCTCTGTT-3'; R, 5'-TGGTGGTAGCGA AGCGTAAG-3'), miR-188-3p (F, 5'-CTCCACATGCAGGG-3'; R, 5'-GTGCAGGGTCCGAGGT-3'), U6 (F, 5'-CTCGCTTCG GCAGCAC-3'; R, 5'-AACGCTTACGAATTTGCGT-3'), and GAPDH (F, 5'-GAATGGGCAGCCGTTAGGAA-3'; R, 5'-AAA AGCATCACCCGGAGGAG-3'). U6 or GAPDH served as a reference control. RNA level was computed using the 2<sup>-ΔΔCt</sup> method (17).

### Western Blotting

Cells were lysed using RNA immunoprecipitation (RIPA) (Solarbio), and protein samples were collected via centrifugation. Proteins of 20 μg were loaded on sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to nitrocellulose membranes (Solarbio). The membranes were blocked in 5% fat-free milk and incubated with primary antibodies anti-TRPC6 (ab62461, Abcam, Cambridge, MA,



**FIGURE 1 |** The levels of circ\_0003204, miR-188-3p, and TRPC6 in ox-LDL-treated HCTAECs and THP-1 cells. Circ\_0003204 expression (A,B) (one-way ANOVA), miR-188-3p expression (C,D) (one-way ANOVA), and TRPC6 protein level (E,F) (one-way ANOVA) were detected in HCTAECs and THP-1 cells after exposure to various doses of ox-LDL for 24 h. \* $p < 0.05$ . TRPC6, transient receptor potential canonical channel 6; ox-LDL, oxidized low-density lipoprotein; HCTAECs, human carotid artery endothelial cells.

USA), anti-BAX (ab104156, Abcam), anti-BCL2 (ab194583, Abcam), or anti- $\beta$ -actin (ab8227, Abcam) and the secondary antibody (ab205718, Abcam). Next, the bands were exposed to enhanced chemiluminescence (ECL) reagent (Solarbio).

## Dual-Luciferase Reporter Analysis and RNA Immunoprecipitation

The wild-type luciferase reporter plasmid circ\_0003204-WT was constructed via cloning the wild-type sequence of circ\_0003204 into psiCHECK-2 vectors (YouBio, Changsha, China). HCTAECs and THP-1 cells were co-transfected with circ\_0003204-WT, circ\_0003204-WT+miR-con, circ\_0003204-WT+miR-188-3p mimic, circ\_0003204-WT+miR-188-3p mimic+pcDNA, or circ\_0003204-WT+miR-188-3p mimic+TRPC6 overexpression vector for 24 h. Next, the luciferase intensity was analyzed via a dual-luciferase analysis kit.

The Magna RIP Kit (Sigma, St. Louis, MO, USA) was exploited for RIP analysis;  $1 \times 10^7$  HCTAECs and THP-1 cells were lysed and interacted with anti-Ago2 or anti-IgG-conjugated magnetic beads for 6 h. The enrichment of circ\_0003204, miR-188-3p, and TRPC6 in the complex was detected by qRT-PCR.

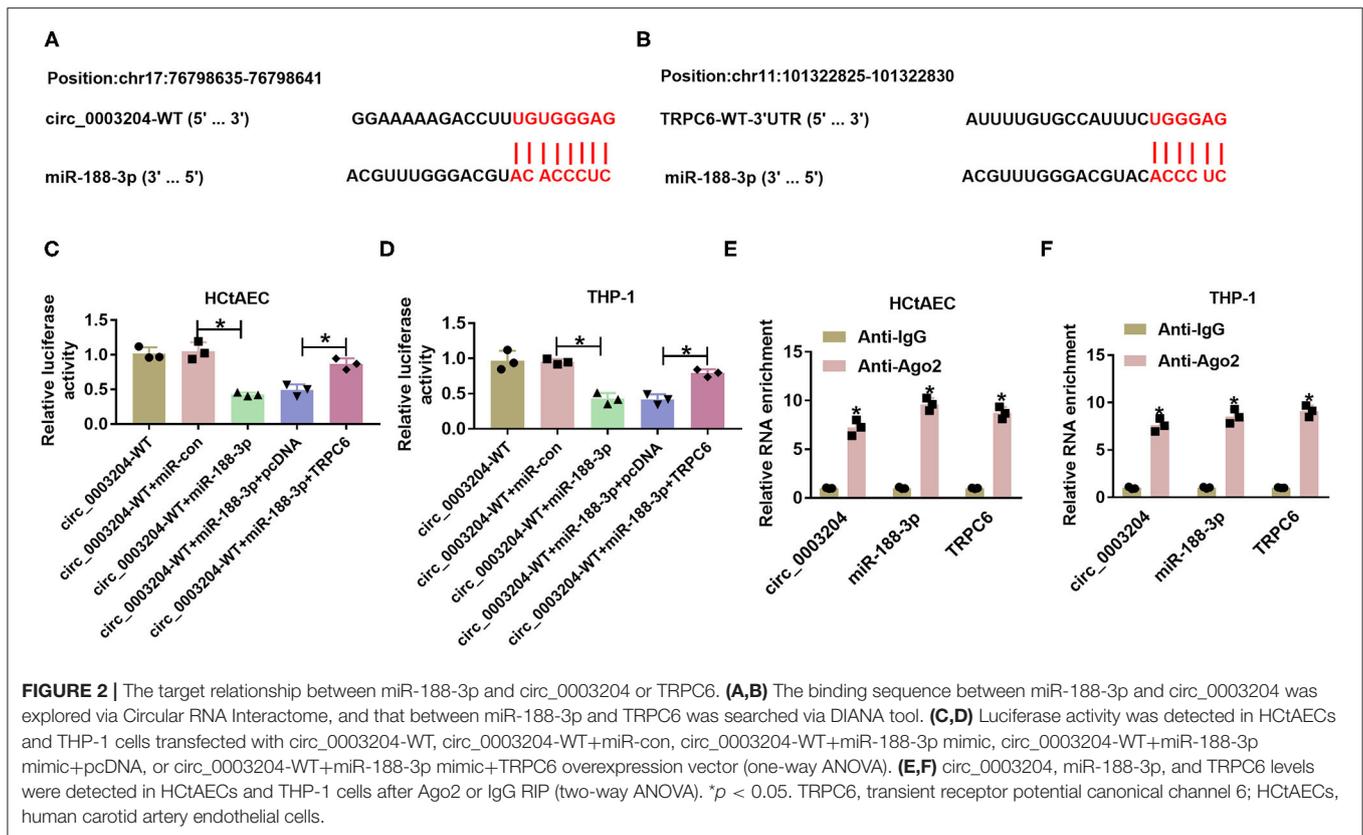
## Cell Transfection

Circ\_0003204 overexpression vector was synthesized by cloning circ\_0003204 sequence into pcDNA3.1 circRNA mini vector,

with the pcDNA3.1 circRNA mini vector (Addgene, Cambridge, MA, USA) as negative control (vector). TRPC6 overexpression vector was generated via inserting the full length of TRPC6 (accession: NM\_004621.6) sequence into pcDNA3.1 vector, with the pcDNA3.1 vector (Addgene) as negative control (pcDNA). SiRNA for circ\_0003204 (si-circ\_0003204, 5'-CCGCAUGGG GCUGUGUACACCU-3'), negative control of siRNA (si-con, 5'-AAGACAUUGUGUGUCCGCCTT-3'), miR-188-3p mimic (5'-CUCCCACAUGCAGGGUUUGCA-3'), negative control of mimic (miR-con, 5'-ACGUGACACGUUCGGAGAATT-3'), miR-188-3p inhibitor (anti-miR-188-3p, 5'-UGCAAACCG ACUUGUGGGAG-3'), and negative control of inhibitors (anti-miR-con, 5'-UGAGCUGCAUAGAGUAGUGAUUA-3') were synthesized via RiboBio (Guangzhou, China). HCTAECs and THP-1 cells were transfected using Lipofectamine 2000 (Thermo Fisher).

## Cell Viability

Cell viability was tested by 3-(4,5-dimethyl-2-thiazolyl)-2,5-diphenyl-2H-tetrazolium bromide (MTT) analysis;  $1 \times 10^4$  HCTAECs were added into 96-well plates overnight and stimulated via 100  $\mu$ g/ml of ox-LDL for 24 h. Next, culture medium was changed to fresh one plus 0.1 mg/ml MTT (Solarbio). After culture for 4 h, the medium was removed, and each well was added with 100  $\mu$ l of dimethyl sulfoxide (DMSO)



(Beyotime, Shanghai, China). The absorbance was examined at 570 nm using a microplate reader.

### 5-Ethynyl-2'-Deoxyuridine Assay

After relevant transfection and ox-LDL treatment, 5-ethynyl-2'-deoxyuridine (EdU) assay kit (Beyotime) was used for cell proliferation. In brief, cells were seeded into 24-well plates ( $5 \times 10^3$  cells/well), and then EdU was added for 2 h of incubation. After that, the cells were fixed with 4% paraformaldehyde (Sigma) and mixed with 0.5% Triton X-100 (Sigma), followed by incubation with Apollo and DAPI. Last, EDU-positive cells were quantified.

### Caspase-3 Activity and Flow Cytometry

For detection of caspase-3 activity,  $4 \times 10^5$  HCTAECs were added into 6-well plates and exposed to 100  $\mu$ g/ml of ox-LDL. Next, cells were lysed for caspase-3 activity analysis using a caspase-3 assay kit (Abcam) according to the instruction of the manufacturer.

For analysis of cell apoptotic rate,  $2 \times 10^5$  HCTAECs were placed into 6-well plates overnight and then stimulated via 100  $\mu$ g/ml of ox-LDL for 24 h. Next, cells were detected using Annexin V-FITC apoptosis detection kit (Sigma). The apoptotic cells were examined using a flow cytometer (Agilent, Hangzhou, China).

### Enzyme-Linked Immunosorbent Assay

The inflammatory response was assessed via analysis of the levels of IL-1 $\beta$ , IL-6, and TNF- $\alpha$ ;  $1 \times 10^5$  THP-1 cells were placed into 12-well plates overnight and then treated via 100  $\mu$ g/ml of

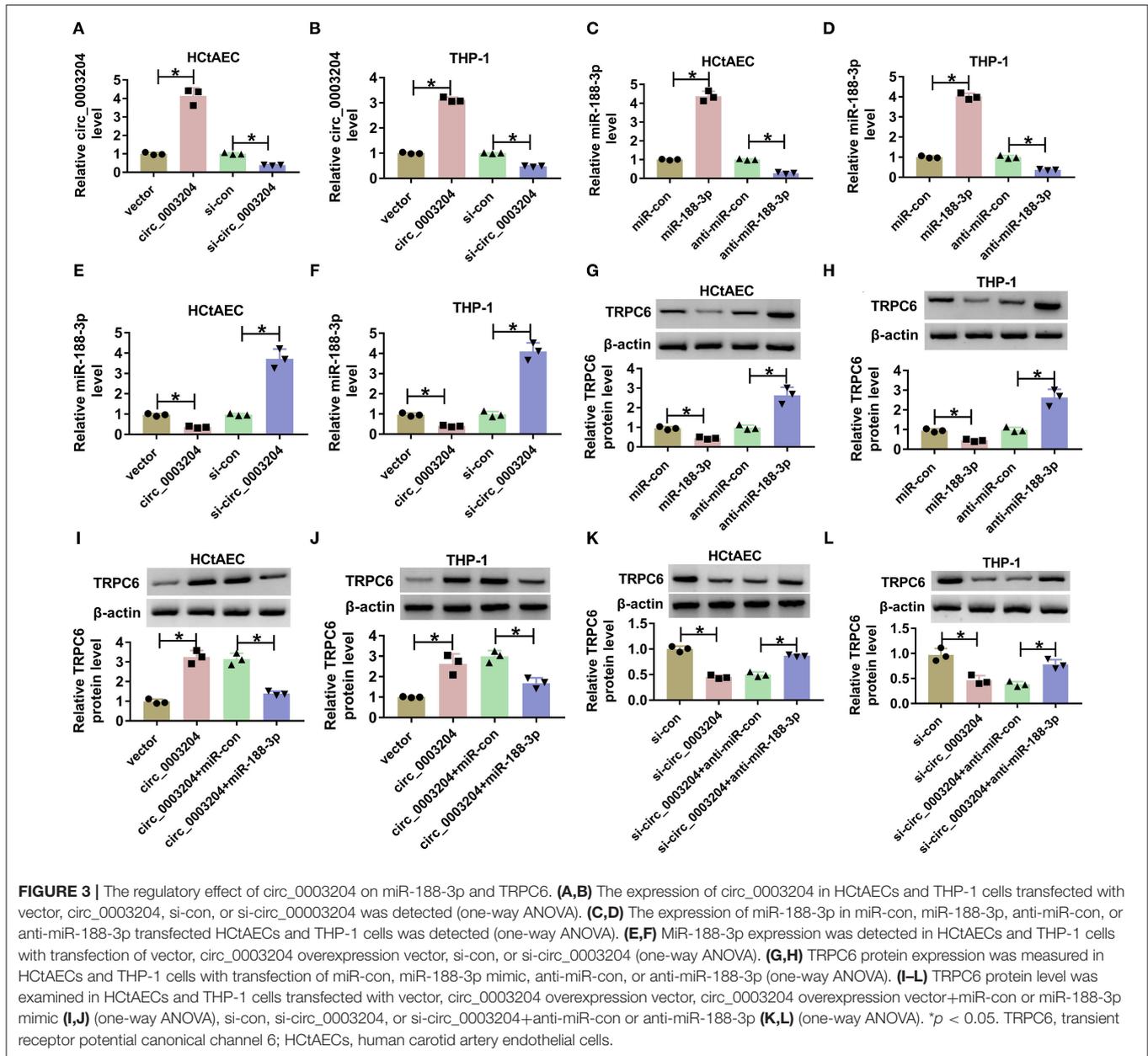
ox-LDL for 24 h. Next, the medium was collected and used for analysis of IL-1 $\beta$ , IL-6, and TNF- $\alpha$  levels using specific ELISA kits (Thermo Fisher) following the instructions of the manufacturer.

### Detection of Malondialdehyde, Superoxide Dismutase, and Reactive Oxygen Species Levels

The oxidative stress was analyzed by evaluation of malondialdehyde (MDA), superoxide dismutase (SOD), and reactive oxygen species (ROS) levels;  $4 \times 10^4$  THP-1 cells were placed into 24-well plates overnight and then challenged via 100  $\mu$ g/ml of ox-LDL for 24 h. Next, cells were lysed; and the lysis solution was used for analysis of MDA and SOD levels using MDA or SOD Assay kit (Sigma) following the instructions of the manufacturer. The levels of MDA and SOD were normalized to total protein in cells. In addition,  $2 \times 10^4$  THP-1 cells were added into 96-well plates overnight and then treated via 100  $\mu$ g/ml of ox-LDL for 24 h. Next, cells were analyzed using a cellular ROS assay kit (Abcam). The ROS level was normalized to the control group (not treated with ox-LDL).

### Statistical Analysis

The experiments were performed 3 times  $\times$  3 replicates. The data were presented as mean  $\pm$  SD. The difference was compared via Student's *t*-test or ANOVA with Dunnett's test using SPSS 20.0. *p* < 0.05 indicated the statistical significance.



## RESULTS

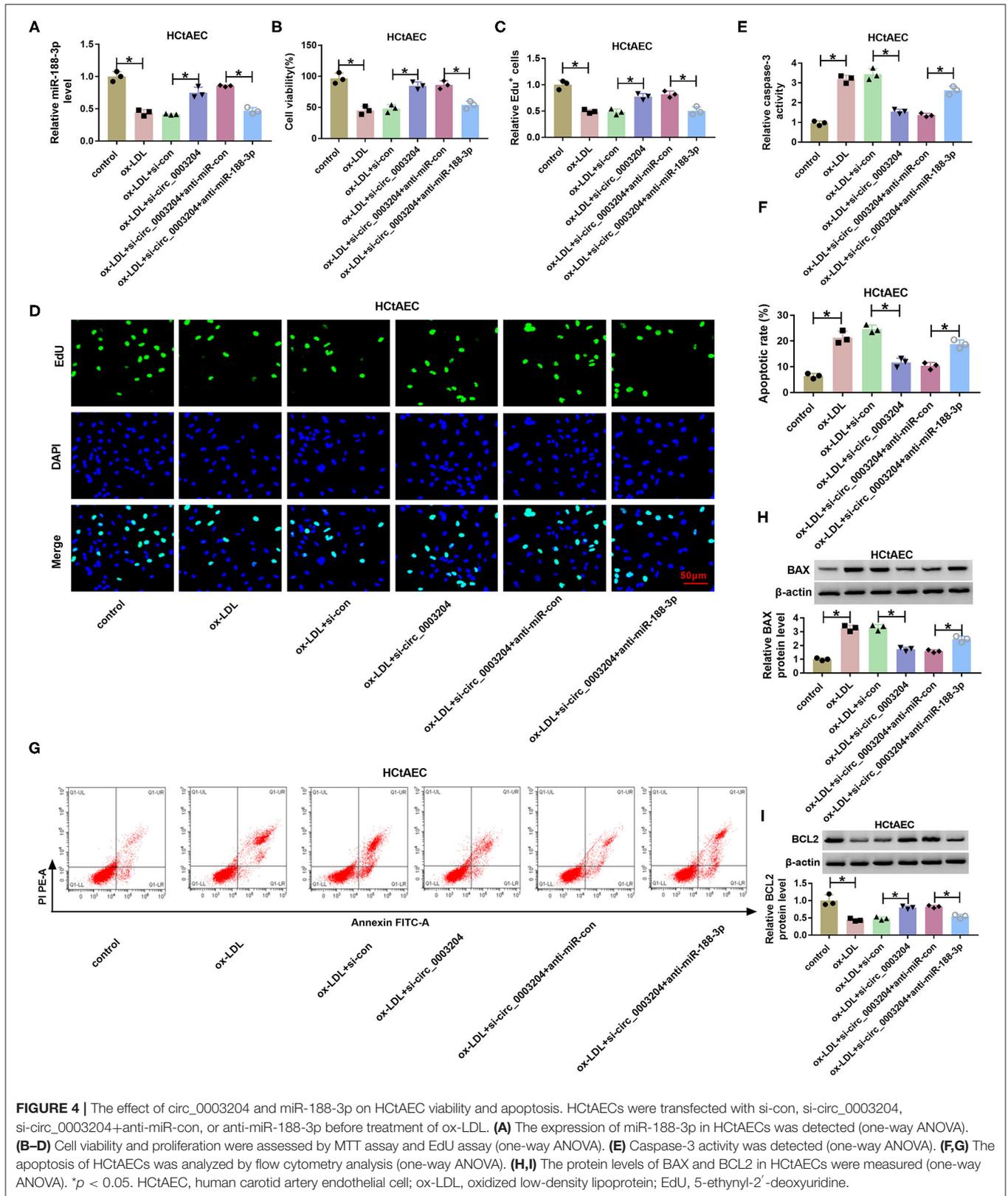
### Circ\_0003204 and Transient Receptor Potential Canonical Channel 6 Levels Are Increased and microRNA-188-3p Expression Is Decreased in Oxidized Low-Density Lipoprotein-Treated Human Carotid Artery Endothelial Cells and THP-1 Cells

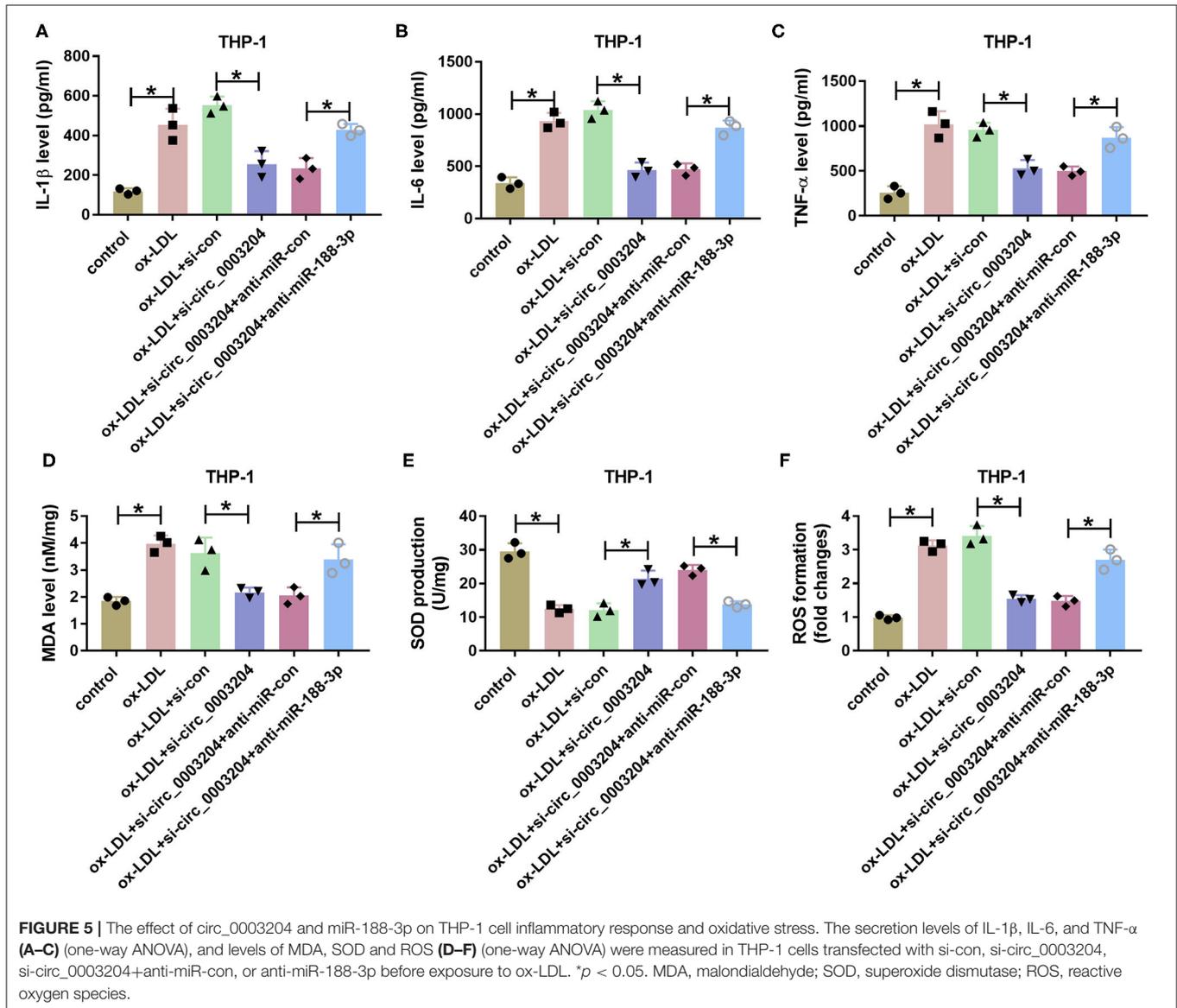
To explore whether circ\_0003204, miR-188-3p, and TRPC are implicated in carotid AS, their levels were detected in ox-LDL-challenged HCTAECs and THP-1 cells. Circ\_0003204 expression was evidently increased in ox-LDL-treated HCTAECs and THP-1 cells in a dose-dependent manner (Figures 1A,B).

Furthermore, miR-188-3p level was progressively reduced as the elevation of ox-LDL (Figures 1C,D). Additionally, TRPC6 protein expression was markedly elevated in the two cell lines after exposure to different doses of ox-LDL (Figures 1E,F). These data indicated that circ\_0003204, miR-188-3p, and TRPC might be associated with ox-LDL-induced injury.

### Circ\_0003204 Regulates Transient Receptor Potential Canonical Channel Expression via Mediating microRNA-188-3p

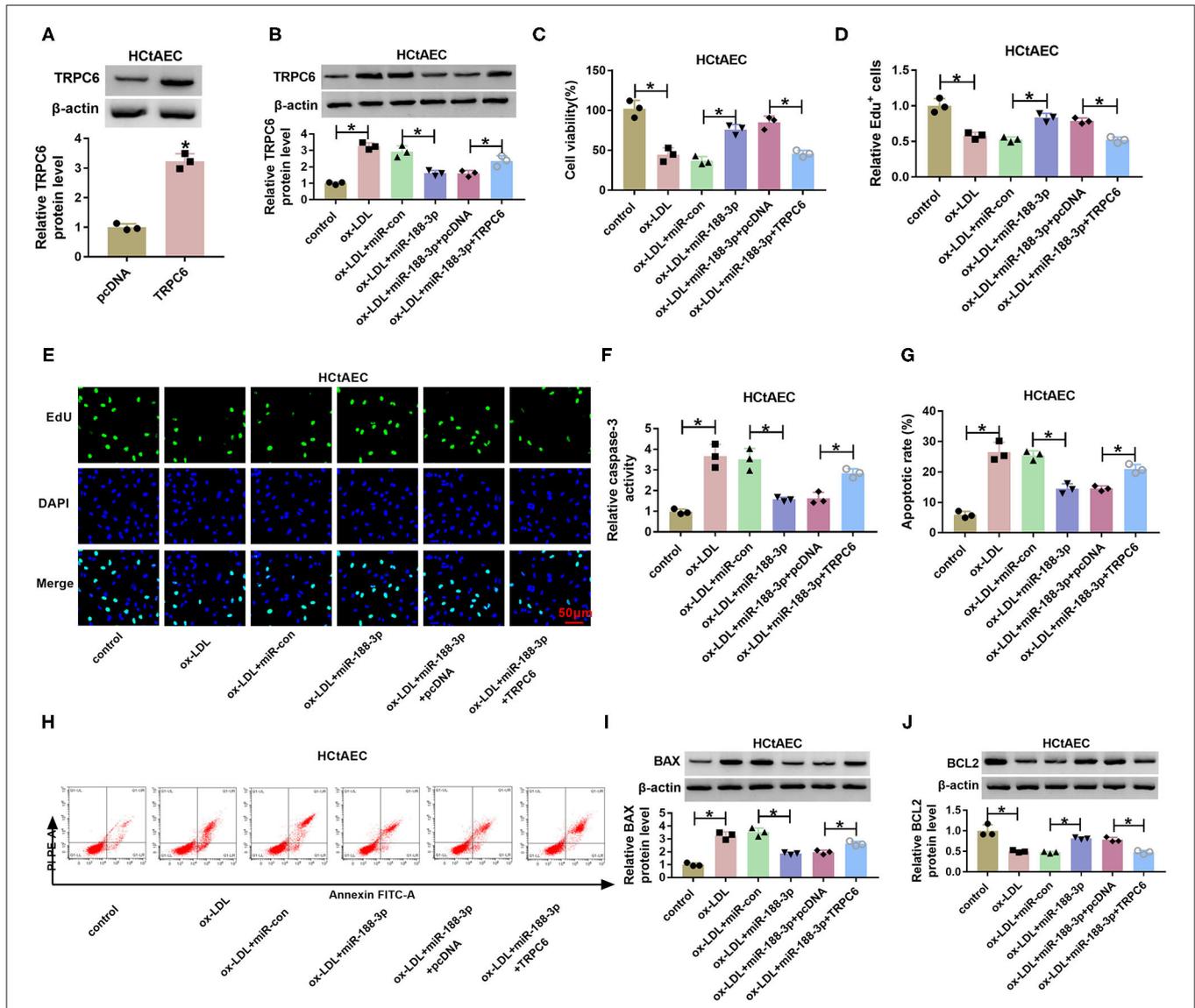
We further analyzed whether circ\_0003204 could act as a ceRNA to regulate the miR-188-3p/TRPC axis. The predicted





binding sequences between miR-188-3p and circ\_0003204 or TRPC are displayed in **Figures 2A,B**. To identify their interactions, the dual-luciferase reporter analysis was performed in HCTaECs and THP-1 cells transfected with circ\_0003204-WT, circ\_0003204-WT+miR-con, circ\_0003204-WT+miR-188-3p mimic, circ\_0003204-WT+miR-188-3p mimic+pcDNA, or TRPC overexpression vector. The luciferase activity of circ\_0003204-WT evidently declined via miR-188-3p overexpression, which was impaired via introduction of TRPC6 (**Figures 2C,D**). Furthermore, the RIP analysis using anti-Ago2 revealed that circ\_0003204, miR-188-3p, and TRPC were enriched in the same complex (**Figures 2E,F**). As exhibited in **Figures 3A,B**, transfection of the circ\_0003204 overexpression vector led to an elevation in circ\_0003204 level, whereas transfection of si-circ\_0003204 led to a reduction in circ\_0003204 level in both HCTaECs and THP-1 cells.

Transfection of MiR-188-3p mimic increased miR-188-3p expression, but introduction of anti-miR-188-3p reduced miR-188-3p expression in both HCTaECs and THP-1 cells (**Figures 3C,D**). Additionally, miR-188-3p expression was significantly reduced via circ\_0003204 overexpression and enhanced via circ\_0003204 knockdown (**Figures 3E,F**). Moreover, TRPC6 protein level was negatively regulated by miR-188-3p in HCTaECs and THP-1 cells (**Figures 3G,H**). As expected, TRPC protein expression was increased by circ\_0003204 overexpression, whereas this elevation was reversed after miR-188-3p overexpression (**Figures 3I,J**). In contrast, circ\_0003204 silencing resulted in a decrease in TRPC protein levels in HCTaECs and THP-1 cells, but this reduction was weakened after miR-188-3p inhibition (**Figures 3K,L**). These data indicated that circ\_0003204 could function as a ceRNA for miR-188-3p to modulate TRPC6.



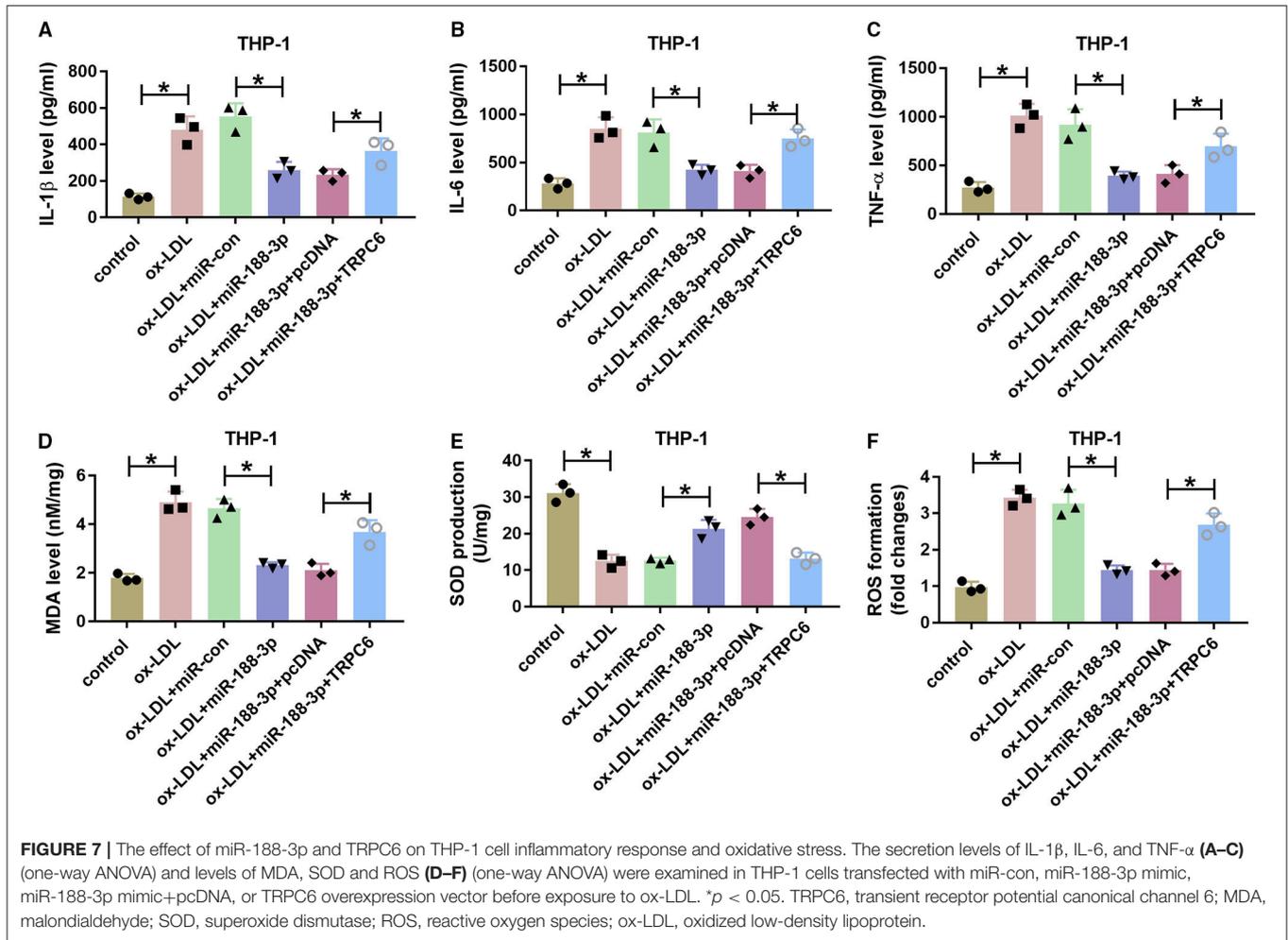
**FIGURE 6 |** The effect of miR-188-3p and TRPC6 on HCTAEC viability and apoptosis. **(A)** TRPC6 protein level was measured (Student's *t*-test). **(B–J)** HCTAECs transfected with miR-con, miR-188-3p mimic, miR-188-3p mimic+pcDNA, or TRPC6 overexpression vector before stimulation of ox-LDL. **(B)** TRPC6 protein level was measured (one-way ANOVA). **(C–E)** Cell viability and cell proliferation were evaluated by MTT and EdU assays (one-way ANOVA). **(F)** Caspase-3 activity was examined (one-way ANOVA). **(G,H)** The apoptosis of HCTAECs was analyzed by flow cytometry analysis (one-way ANOVA). **(I,J)** The protein levels of BAX and BCL2 were measured (one-way ANOVA). \**p* < 0.05. HCTAEC, human carotid artery endothelial cell; TRPC6, transient receptor potential canonical channel 6; ox-LDL, oxidized low-density lipoprotein; EdU, 5-ethynyl-2'-deoxyuridine.

### Circ\_0003204 Knockdown Attenuates Oxidized Low-Density Lipoprotein-Induced Injury via Regulating microRNA-188-3p in Human Carotid Artery Endothelial Cells and THP-1 Cells

To probe into whether circ\_0003204 mediated ox-LDL-induced HCTAEC injury and through miR-188-3p, HCTAECs were transfected with si-con, si-circ\_0003204, si-circ\_0003204+anti-miR-con, or anti-miR-188-3p before treatment of ox-LDL. qRT-PCR assay showed that circ\_0003204 knockdown increased miR-188-3p level in ox-LDL-induced

HCTAECs, while introduction of anti-miR-188-3p reversed the effect (Figure 4A). As displayed in Figures 4B–D, circ\_0003204 knockdown mitigated ox-LDL-induced viability and proliferation inhibition in HCTAECs, which was weakened by miR-188-3p inhibition. Moreover, circ\_0003204 silencing weakened ox-LDL-induced apoptosis by decreasing caspase-3 activity and regulating BAX and BCL2 protein levels in HCTAECs, which was abolished via miR-188-3p downregulation (Figures 4E–I).

In addition, the effect of circ\_0003204 on ox-LDL-induced THP-1 cell damage was assessed. As described in Figures 5A–C, interference of circ\_0003204 alleviated ox-LDL-caused



inflammatory response via decreasing IL-1 $\beta$ , IL-6, and TNF- $\alpha$ , which was reversed via miR-188-3p knockdown. Besides, circ\_0003204 silencing attenuated ox-LDL-induced oxidative stress via decreasing MDA and ROS levels and increasing SOD level, and these events were weakened by miR-188-3p knockdown (Figures 5D–F). These results suggested that circ\_0003204 knockdown weakened ox-LDL-induced HCtAEC and THP-1 cell injury via mediating miR-188-3p.

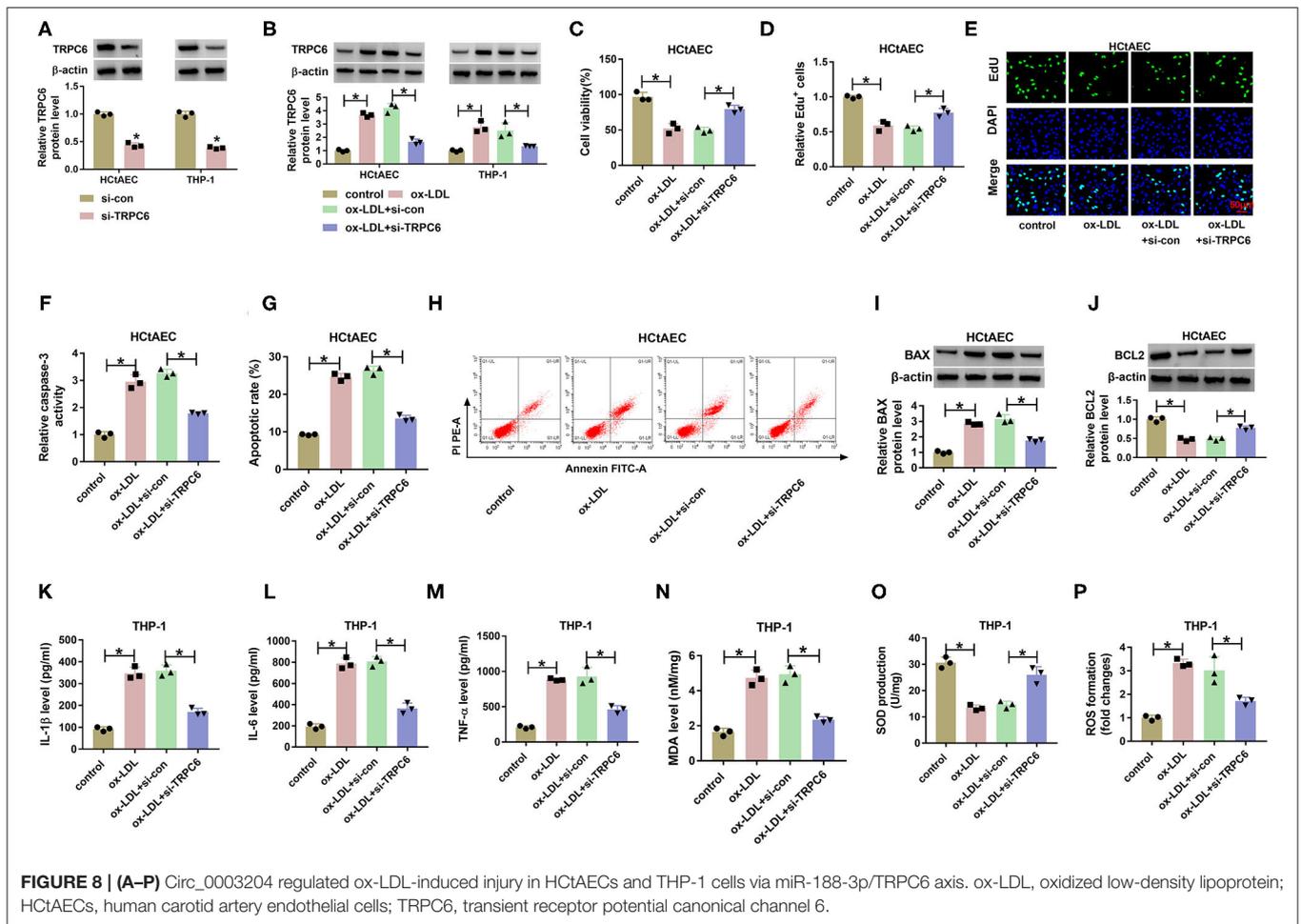
### Transient Receptor Potential Canonical Channel 6 Knockdown Weakens Oxidized Low-Density Lipoprotein-Induced Injury in Human Carotid Artery Endothelial Cells and THP-1 Cells

To analyze the effects of TRPC6 knockdown on ox-LDL-induced injury in HCtAECs and THP-1 cells, we knocked out TRPC6 by transfection with si-TRPC6. The transfection efficiency is exhibited in Supplementary Figure 1A. Moreover, introduction of si-TRPC6 weakened the elevation of TRPC6 mediated by ox-LDL stimulation (Supplementary Figure 1B).

As expected, TRPC6 downregulation weakened ox-LDL-induced viability and proliferation inhibition (Supplementary Figures 1C–E). Moreover, the elevated caspase-3 activity, apoptotic rate, and BAX protein levels and the decreased BCL-2 protein levels in HCtAECs and THP-1 cells caused by ox-LDL were mitigated by TRPC6 knockdown (Supplementary Figures 1F–I). In addition, ox-LDL-induced cell inflammation (Supplementary Figures 1K–M) and oxidative stress (Supplementary Figures 1N–P) were impaired after TRPC6 silencing. These results suggested that TRPC6 silencing weakens ox-LDL-induced injury in HCtAECs and THP-1 cells.

### MicroRNA-188-3p Overexpression Mitigates Oxidized Low-Density Lipoprotein-Induced Injury via Mediating Transient Receptor Potential Canonical Channel 6 in Human Carotid Artery Endothelial Cells and THP-1 Cells

As exhibited in Figure 6A, transfection of the TRPC6 overexpression vector increased TRPC6 protein level in



HCTaECs. To test whether miR-188-3p was associated with ox-LDL-induced HCTaEC injury through TRPC6, HCTaECs were transfected with miR-con, miR-188-3p mimic, miR-188-3p mimic+pcDNA, or TRPC6 overexpression vector before exposure to ox-LDL. As exhibited in **Figures 6B–E**, miR-188-3p overexpression mitigated ox-LDL-induced viability and proliferation suppression in HCTaECs, which was abrogated after TRPC6 overexpression. Furthermore, overexpression of miR-188-3p mitigated ox-LDL-caused apoptosis via reducing caspase-3 and modulating BAX and BCL2 protein levels, which was reversed by TRPC6 upregulation (**Figures 6F–J**).

Additionally, the effect of miR-188-3p on ox-LDL-induced damage in THP-1 cells was tested. As shown in **Figures 7A–C**, overexpression of miR-188-3p alleviated ox-LDL-induced inflammatory response via reducing the levels of IL-1 $\beta$ , IL-6, and TNF- $\alpha$  in THP-1 cells, which were overturned by TRPC6 overexpression. Besides, miR-188-3p overexpression attenuated ox-LDL-induced oxidative stress by inhibiting MDA and ROS levels and elevating SOD level, while these events were reversed after TRPC6 restoration (**Figures 7D–F**). These data indicated that miR-188-3p overexpression attenuated ox-LDL-induced damage in HCTaECs and THP-1 cells via targeting TRPC6.

## DISCUSSION

AS causes high morbidity and mortality all around the world (18). Carotid artery AS is a common form of AS (19). The inflammatory response, oxidative stress, and endothelial cell apoptosis are implicated in AS progression (20–22). CircRNAs play a key role in the development of AS (23). This research focused on the function and mechanism of circ\_0003204 in the regulation of ox-LDL-induced inflammatory response, oxidative stress, and endothelial cell apoptosis. Here, we first found that circ\_0003204 knockdown could mitigate ox-LDL-induced injury in HCTaECs and THP-1 cells.

We established ox-LDL-stimulated HCTaECs and THP-1 cells, and we found that circ\_0003204 expression was elevated in the two cell lines, which was consistent with that in HAECs or HUVECs (8, 9). Hence, we assumed the increase in circ\_0003204 induced by abnormal ox-LDL might be correlated with carotid artery AS development. Previous studies suggested that ox-LDL could induce endothelial cell injury and that ox-LDL-stimulated HCTaECs could be used to assess the pathogenesis of carotid artery AS *in vitro* (15, 24, 25). Similarly, our study also found that ox-LDL caused HCTaEC viability inhibition and apoptosis promotion and that circ\_0003204 knockdown

weakened ox-LDL-exposed HCtAEC injury. In addition, ox-LDL-mediated THP-1 cell damage is also involved in the development of carotid artery AS (15, 26, 27). By detecting the pro-inflammatory cytokine levels and oxidative stress-related markers, we confirmed that circ\_0003204 silencing attenuated ox-LDL-induced inflammatory response and oxidative stress in THP-1 cells. Collectively, inhibition of circ\_0003204 played an inhibiting effect in ox-LDL-induced injury in AS.

Previous studies suggested that circRNA-mediated ceRNA network is the important mechanism in cardiovascular diseases and ox-LDL-induced injury (28, 29). The former work has confirmed that circ\_0003204 could act as a ceRNA to regulate the miR-370-3p/TGF $\beta$ 2 axis (8). Here, we first explored the potential association among circ\_0003204, miR-188-3p, and TRPC6. In addition, we found that circ\_0003204 could positively regulate TRPC6 expression by binding to miR-188-3p, indicating that circ\_0003204 might function as a ceRNA for miR-188-3p to mediate TRPC6. Our study found that miR-188-3p overexpression weakened ox-LDL-induced viability inhibition and apoptosis promotion in HCtAECs. Furthermore, we confirmed miR-188-3p weakened ox-LDL-induced inflammatory response in THP-1 cells, which was similar to that in a previous study (12). Besides, our results also displayed that miR-188-3p could mitigate ox-LDL-induced oxidative stress in THP-1 cells. These data indicated the protective function of miR-188-3p in ox-LDL-induced damage. Additionally, miR-188-3p knockdown reversed the effect of circ\_0003204 silencing on ox-LDL-induced damage in the two cell lines, implying that circ\_0003204 could regulate ox-LDL-induced cell injury by mediating miR-188-3p in HCtAECs and THP-1 cells.

Next, we validated the targeting interaction between miR-188-3p and TRPC6. Negri et al. and Thilo et al. reported that TRPC6 was abnormally expressed in vascular endothelial cells, and its elevation was associated with cardiovascular disease development (30, 31). Moreover, Zhang et al. suggested that TRPC6 could contribute to AS development by promoting endothelial cell apoptosis (14). Similarly, our study also confirmed that the inhibitive role of miR-188-3p on ox-LDL-mediated cell injury in HCtAECs and THP-1 cells was partly reversed by TRPC6 overexpression. Collectively, we concluded that circ\_0003204 could regulate ox-LDL-induced cell injury by mediating miR-188-3p and TRPC6. However, it did not indicate the biological role of circ\_0003204 in carotid artery AS *in vivo* because of the alteration of microenvironment. The animal models have

been widely used to assess the pathogenesis of AS *in vivo* (32). Therefore, the preclinical experiments using animals would be performed to test the function and mechanism of circ\_0003204 in carotid artery AS in future.

In conclusion, circ\_0003204 knockdown could attenuate ox-LDL-induced HCtAEC apoptosis and THP-1 cell inflammatory response and oxidative stress via modulating miR-188-3p/TRPC6 axis in a ceRNA network (Figure 8). This study indicated that this ceRNA crosstalk might be associated with ox-LDL-induced injury in carotid artery AS and that circ\_0003204 might act as a target for carotid artery AS therapy.

## DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/Supplementary Material, further inquiries can be directed to the corresponding author/s.

## AUTHOR CONTRIBUTIONS

WP designed and performed the research and wrote the manuscript. SL, SC, JY, and ZS analyzed the data. All authors have read and approved the final manuscript.

## FUNDING

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## SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fcvm.2021.731890/full#supplementary-material>

**Supplementary Figure 1** | TRPC6 knockdown weakens ox-LDL-induced injury in HCtAEC and THP-1 cells. **(A)** Western blot showing TRPC6 protein levels in HCtAEC and THP-1 cells transfected with si-con or si-TRPC6 (Student's *t*-test). **(B–P)** HCtAEC cells were transfected with si-con or si-TRPC6 and then treated with ox-LDL. **(B)** Western blot showing TRPC6 protein levels in HCtAEC and THP-1 cells (one-way ANOVA). **(C–E)** MTT and EdU assays (one-way ANOVA) were performed to analyze cell viability and proliferation (one-way ANOVA). **(F)** Evaluation of Caspase-3 activity was performed (one-way ANOVA). **(G,H)** Flow cytometry assays were carried out to analyze cell apoptosis (one-way ANOVA). **(I,J)** The protein levels of BAX and BCL2 in HCtAEC cells were measured (one-way ANOVA). **(K–P)** Measurements of IL-1 $\beta$ , IL-6, TNF- $\alpha$ , MDA, SOD, and ROS were conducted (one-way ANOVA). \**P* < 0.05.

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