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## XJB-5-131 protects chondrocytes from ferroptosis to alleviate osteoarthritis progression via restoring Pebp1 expression<sup>☆</sup>

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## ABSTRACT

**Background:** Osteoarthritis (OA) is the most common age-related musculoskeletal disease. However, there is still a lack of therapy that can modify OA progression due to the complex pathogenic mechanisms. The aim of the study was to explore the role and mechanism of XJB-5-131 inhibiting chondrocytes ferroptosis to alleviate OA progression.

**Methods:** We treated tert-butyl hydroperoxide (TBHP)-induced ferroptosis of mouse primary chondrocytes with XJB-5-131 in vitro. The intracellular ferroptotic hallmarks, cartilage anabolic and catabolic markers, ferroptosis regulatory genes and proteins were detected. Then we established a mouse OA model via destabilization of the medial meniscus (DMM) surgery. The OA mice were treated with intra-articular injection of XJB-5-131 regularly (2 μM, 3 times per week). After 4 and 8 weeks, we performed micro-CT and histological examination to evaluate the protection role of XJB-5-131 in mouse OA subjects. RNA sequencing analysis was performed to unveil the key downstream gene of XJB-5-131 exerting the anti-ferroptotic effect in OA.

**Results:** XJB-5-131 significantly suppressed TBHP-induced increases of ferroptotic hallmarks (ROS, lipid peroxidation, and Fe<sup>2+</sup> accumulation), ferroptotic drivers (Ptgs2, Pgd, Tfrc, Atf3, Cdo1), while restored the expression of ferroptotic suppressors (Gpx4, Fth1). XJB-5-131 evidently promoted the expression of cartilage anabolic and decreased the expression of cartilage catabolic markers. Moreover, intra-articular injection of XJB-5-131 significantly inhibited the expression of Cox2 and Mmp13, while promoted the expression of Col2a1, Gpx4 and Fth1 in DMM-induced mouse articular cartilage. Further, we identified Pebp1 as a potential target of XJB-5-131 by RNA sequencing analysis. The anti-ferroptosis and chondroprotective effects of XJB-5-131 were significantly diminished by Locostatin, a specific antagonist of Pebp1.

<sup>☆</sup> All authors have read and approved the manuscript and agreed to be accountable for all aspects of the work.

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**Conclusion:** XJB-5-131 significantly protects chondrocytes from ferroptosis in TBHP-induced mouse primary chondrocytes and DMM surgery-induced OA mice model via restoring the expression of Pebp1. XJB-5-131 is a potential therapeutic drug in the management of OA progression.

## The translational potential of this article

We demonstrate that XJB-5-131 is a potential therapeutic drug in the management of OA, which provides a new breakthrough in the mechanism treatment of OA with clinical potential.

### 1. Introduction

Osteoarthritis (OA) is the most prevalent age-related musculoskeletal disease characterized by swelling and pain of involved joints, which can be accompanied by joint deformity, movement limitation, and even disability in severe cases [1–4]. Although OA has been recognized for a long time, however, there is still a lack of therapy modifying its progression due to the complex pathogenic mechanisms. The U.S. Food and Drug Administration (FDA) classified OA as “a serious disease with unmet medical need” [5]. There was an urgent need to develop new drugs for the treatment of OA.

Cartilage degeneration has been recognized as the main pathogenesis of OA progression, in which injury of chondrocytes, the only one cell type in cartilage, directly accelerated this process [6–8]. Multiple types of chondrocyte death have been identified in cartilage degeneration [8, 9]. Recently, chondrocyte ferroptosis has been suggested to be directly involved in OA cartilage degeneration [10]. Ferroptosis is a novel form of iron-dependent programmed cell death characterized by lipid peroxidation, accumulated ferrous ions, and depleted glutathione, which is significantly different from previously identified programmed cell death at the genetic, morphological, and biological levels [11–13]. Increasing studies focused on chondrocyte ferroptosis in OA [14–17], whereas, few investigations have been reported to inhibit chondrocyte ferroptosis and alleviate OA. The nitroxide XJB-5-131 is an antioxidant targeting mitochondria and has achieved satisfactory efficacy in a variety of diseases [18–23], as well as being considered a potential inhibitor of ferroptosis [24]. However, the role and mechanism of XJB-5-131 in OA progression have not been investigated.

In the current study, we explored the anti-ferroptotic effects of XJB-5-131 with primary mouse chondrocytes and mouse OA model induced by destabilization of the medial meniscus (DMM) surgery. We found that XJB-5-131 exhibited anti-ferroptotic effect of chondrocyte and alleviated OA progression. These results have important implications for the development of novel therapeutic agents based on the pathogenesis of OA.

### 2. Materials and methods

#### 2.1. Ethics statement

All experimental procedures were in accordance with the Declaration of Helsinki. The animal experimental procedures were performed in strict accordance with the guidelines of the Animal Care and Use Committee of Nanjing Drum Tower Hospital (2020AE01102).

#### 2.2. Isolation and culture of mouse primary chondrocytes

The murine primary chondrocytes were isolated and cultured referring to previous protocols [25]. Briefly, the cartilage tissues from the 3-day-old C57BL/6 mice (purchased from Model Animal Research Center of Nanjing University) were digested in 0.2 % type II collagenase (Gibco, USA) at 37 °C for 4 h. After removing soft tissues and filtrating with a cell strainer (Corning, USA), the isolated chondrocytes were cultured in Dulbecco's modified Eagle's medium contained 1 g/L

glucose (DMEM/F12) (Gibco, USA) supplemented with 1 % penicillin-streptomycin (Invitrogen, USA) and 10 % fetal bovine serum (Gibco, USA) in the incubator (5 % CO<sub>2</sub>, 37 °C) (Thermo Scientific, USA). The cell culture medium was refreshed every 48 h, and chondrocytes from passages 1 to 3 were used in cell experiments.

#### 2.3. Cell viability assay

To assess the cytotoxicity of tert-butyl hydroperoxide (TBHP) (#MKCH9944, Sigma, USA) to chondrocytes in the presence or absence of XJB-5-131 (#SML2982, Sigma, USA), CCK-8 assay was performed (#CK04, Dojindo, Japan) according to the manufacturer's instruction. Uniformly chondrocytes were seeded in 96-well plates and cultured with TBHP and/or XJB-5-131 for 24 h. After washed by PBS, the cells were then incubated with DMEM/F12 containing 10 % CCK-8 solution for 90min at 37 °C. The OD values were measured at 450 nm with a microplate reader (Thermo Scientific, USA).

#### 2.4. Quantitative real-time PCR

The cellular mRNA isolation and qPCR were performed essentially using RNA-quick Purification Kit (#RN001, ES Science, Shanghai, China), HiScript-TS 5'/3' RACE Kit (RA101, Vazyme Biotech Co.,Ltd, China) and ChamQ™ SYBR Color qPCR Master Mix (Q411, Vazyme, China) according to previous protocols [26]. The primer sequences used were as follows:

Gapdh: 5' – AGGTCGGTGTGAACGGATTG- 3' (forward)  
and 5' – TGTAGACCATGTAGTTGAGGTCA- 3' (reverse)  
Atf3: 5' – GAGGATTTTGCTAACCTGACACC- 3' (forward)  
and 5' – TTGACGGTAACTGACTCCAGC- 3' (reverse)  
Gpx4: 5' – TGTGCATCCCGCGATGATT- 3' (forward)  
and 5' – CCCTGTACTTATCCAGGCAGA- 3' (reverse)  
Pgd: 5' – TGAAGGGTCCTAAGGTGGTCC- 3' (forward)  
and 5' – CCGCCATAATTGAGGGTCCAG- 3' (reverse)  
Cdo1: 5' – GGGGACGAAGTCAACGTGG- 3' (forward)  
and 5' – ACCCCAGCACAGAATCATCAG- 3' (reverse)  
Ptgs2: 5' – TGAGCAACTATTCCAAACCAGC- 3' (forward)  
and 5' – GCACGTAGTCTTCGATCACTATC- 3' (reverse)  
Col2a1: 5' – CCAGATGACCTTCTACGCC- 3' (forward)  
and 5' – TTCAGGGCAGTGTACGTGAAC- 3' (reverse)  
Acan: 5' – GTGCCTATCAGACAAGGTCT- 3' (forward)  
and 5' – GATGCCTTTCACCACGACTTC- 3' (reverse)  
Mmp13: 5' – TCTTTATGGTCCAGGCGATGA- 3' (forward)  
and 5' – ATCAAGGGATAGGGCTGGGT- 3' (reverse)  
Mmp3: 5' – ATGGGCCTGGAACAGTCTTG- 3' (forward)  
and 5' – GTGGGAGTTCCATAGAGGGAC- 3' (reverse)  
Tfrc: 5' – GTTCTGCCAGCCCTTATTAT- 3' (forward)  
and 5' – GCAAGGAAAGGATATGCAGCA- 3' (reverse)

#### 2.5. Intracellular Fe<sup>2+</sup>, ROS and lipid peroxidation detection

Intracellular levels of ferrous iron (Fe<sup>2+</sup>), ROS and lipid peroxidation were measured with Flow Cytometry as previously described [15]. Briefly, chondrocytes were incubated with 1 μM Fe<sup>2+</sup> probe FerroOrange (#F374, Dojindo, Japan), 10 μM ROS probe DCFH-DA (#S0033S, beyotime, China) and 5 μM lipid ROS probe C11-BODIPY (#GC40165, GLPBO, USA) for 40 min in the incubator (37 °C, 5 % CO<sub>2</sub>), respectively. After washing the probes with PBS, the chondrocytes were treated with 50 μM TBHP and/or 2 μM XJB-5-131, 10 μM Locostatin (#A16079, Adooq, China) for 4 h. Then the fluorescent intensity of

chondrocytes was measured by Flow Cytometry (BD Biosciences, USA). FlowJo software (Version 10, USA) was used to analyze the data.

## 2.6. Protein extraction and western blot

The total protein of chondrocytes with different treatment were extracted using a whole Cell Lysis Assay Kit (#KGA1202, Keygen, China). Protein concentration measurements and western blot were performed according to the previously described protocols [26]. The primary antibodies used were as follows: anti-Collage II (#BA0533, Boster, China), anti-MMP13 ((#ab219620, Abcam, UK), anti-Cox2 (#66351-1-Ig, Proteintech, China), anti-FTH1 (#4393, Cell Signaling Technology, USA), anti-GPX4 (#ab125066, Abcam, UK), anti-PEBP1 (#GB113118, Servicebio, China). After HRP-conjugated goat anti-rabbit/mouse secondary antibodies (#BL003A or #BL001A, Biosharp, China) incubation, an ECL Western Blot Kit (Tanon, China) was used to detect signals. Finally, we quantified the protein bands with Image J software (version 1.8.0, USA).

## 2.7. Animals study

Twelve weeks old wild-type C57BL/6 male mice were purchased from the Model Animal Research Center of Nanjing University, and acclimated in specific pathogenfree (SPF) condition. The mice were randomly divided into the following 4 groups: (1) Sham operation (Sham); (2) Sham operation with XJB-5-131 treatment (XJB); (3) Destabilization of medial meniscus (DMM); and (4) DMM with XJB-5-131 treatment (DMM + XJB). The XJB-5-131 (2  $\mu$ M, 8  $\mu$ l, 3 times/week) was intra-articularly injected into the right knees since 1 week after DMM surgery for 4 W or 8 W. The equal volume of saline was injected in the Sham and DMM groups. The DMM procedure were performed as previously described [26].

## 2.8. Micro-CT analysis

Mouse knee joints were fixed in 4 % paraformaldehyde (PFA), and then were evaluated with a micro-CT scanner (mCT80, Scanco Medical AG, Switzerland). As previously described. We analyzed the images and reconstructed the 3D knee joints with Scanco Medical software. The number of osteophytes, subchondral plate thickness (SBP), bone volume (BV), subchondral bone mineral density (BMD), bone volume/tissue volume (BV/TV), trabecular thickness (Tb.Th), trabecular separation (Tb.Sp), trabecular number (Tb.N) were acquired from the Scanco Medical software.

## 2.9. Histological analysis

After treatment, the experimental knee joints were surgically removed and fixed in 4 % PFA, decalcified with 10 % EDTA and embedded in paraffin. The subjects were sliced into continuous coronal 5  $\mu$ m slides by a microtome (Thermo, USA) and stained with safranin O/fast green (Solarbio, China). The Osteoarthritis Research Society International (OARSI) grading system was used to evaluate the cartilage destruction. We also analyzed the thickness of cartilage and the number of chondrocytes.

## 2.10. Immunofluorescence staining

After dewaxing, the histological sections of mouse knee joints were blocked with 5 % bovine serum albumin (BSA) for 1 h at room temperature, then incubated overnight at 4 °C with primary antibodies as follows: anti-Collage II (#BA0533, Boster, China), anti-MMP13 (#ab219620, Abcam, UK), anti-Cox2 (#ab219620, Abcam, UK). Then, the slides were washed with TBST and incubated with a fluorescein-conjugated secondary antibody (#BL003A or #BL001A, Abcam, USA) for 60 min at room temperature. Then we stained the nuclei with DAPI

(#SC3598, Santa Cruz, USA). A fluorescence microscope (Zeiss, Germany) was used to observe and acquire the images.

## 2.11. RNA sequence analysis

Total RNA was extracted from primary chondrocytes after the treatment of TBHP with or without XJB-5-131 for 6 h (n = 3). The RNA was submitted to GeneChem company (Shanghai, China) for RNA sequencing analysis. The Pearson's correlation analysis and heatmaps were performed. In this study, differentially expressed genes (DEGs) were defined as fold changes >2 and P < 0.05. We used Gene ontology (GO) and Kyoto Encyclopedia of Genes and Genomes (KEGG) analysis to assess the biological function of DEGs.

## 2.12. Statistical analysis

The data were expressed as the mean  $\pm$  standard deviation (S.D.). The data were statistically analyzed by GraphPad Prism software (version 8.0, USA) and SPSS software (version 24.0, USA). The quantitative data used represent the results of no less than three independent repeated experiments. Levene method and Shapiro–Wilk test were used for the estimation of data homogeneity of variance and normal distribution, respectively. The data between two groups were compared with unpaired two-tailed t-test. When more than 2 groups, we performed One-way ANOVA statistical analysis. The statistically significant was considered as P < 0.05.

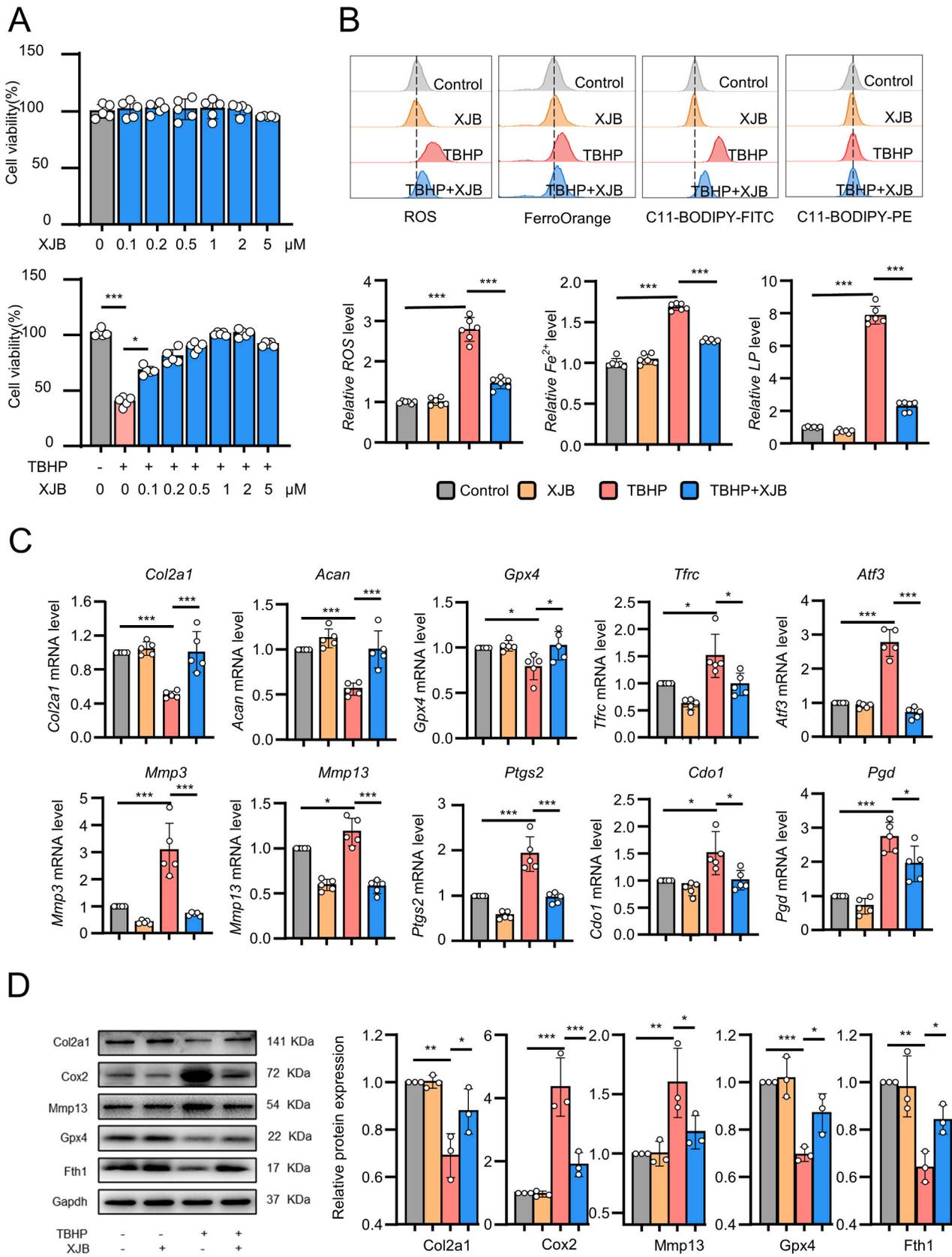
## 3. Results

### 3.1. XJB-5-131 suppresses TBHP-induced chondrocytes ferroptosis in vitro

The cytotoxicity of XJB-5-131 were performed by CCK-8 kit, the results showed that XJB-5-131 at concentrations of 0.1–5  $\mu$ M exhibited no cytotoxic effects on chondrocyte viability while reduced chondrocyte viability but had no significant difference at 5  $\mu$ M concentration. As previously described, we used tert-butyl hydroperoxide (TBHP) (50  $\mu$ M) to induce mouse primary chondrocyte ferroptosis in vitro [15]. The viability of chondrocytes was significantly decreased after 12 h incubation of TBHP, however, the cytotoxicity was significantly alleviated when the concentration of XJB-5-131 increased (Fig. 1A). Next, we found that XJB-5-131 significantly alleviated the ferroptotic hallmarks (intracellular ROS level, Fe<sup>2+</sup> accumulation and lipid peroxidation level) in TBHP-induced chondrocytes by flow cytometry (Fig. 1B). Q-PCR verified that XJB-5-131 promoted the expression of chondrocyte anabolic genes (Col2a1 and Acan) and inhibited chondrocyte catabolic genes (Mmp13 and Mmp3) (Fig. 1C). In addition, under the treatment of TBHP, the ferroptotic suppressor (Gpx4) was evidently up-regulated and ferroptotic drivers (Tfrc, Atf3, Ptg2, Cdo1, Pgd) were significantly down-regulated by XJB-5-131 (Fig. 1C). Moreover, western blot analysis showed that XJB-5-131 significantly promoted the expression of chondrocyte anabolic protein (Col2a1) and ferroptotic suppressors (Gpx4 and Fth1), while markedly decreased chondrocyte metabolic protein (Mmp13) and ferroptotic driver (Cox2) (Fig. 1D). These results suggest that XJB-5-131 plays an anti-ferroptotic role in chondrocytes.

### 3.2. XJB-5-131 alleviates DMM-induced osteophyte formation and subchondral bone sclerosis

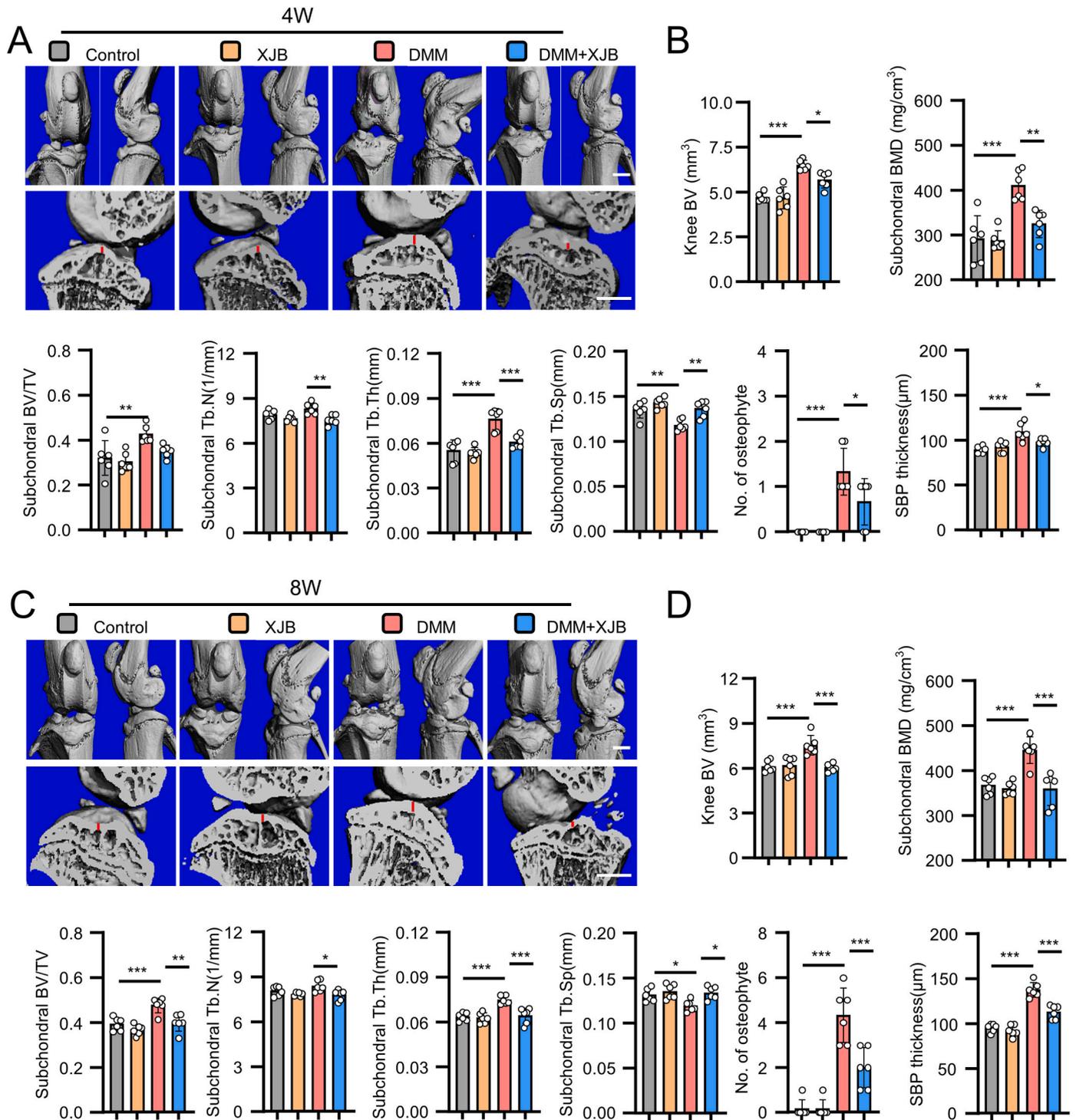
To gain insight into whether XJB-5-131 can suppress chondrocyte ferroptosis and alleviate OA in vivo, we performed intra-articular injection of XJB-5-131 in the OA mouse model. Previous study suggested that OA cartilage degeneration was highly associated with bone redundancy formation and subchondral bone microarchitectural changes [27]. We further analyzed the effect of XJB-5-131 on the bone quality of the mice knee joint. We found osteophytes, subchondral bone



**Fig. 1.** XJB-5-131 suppresses TBHP-induced chondrocytes ferroptosis in vitro. (A) The viability of primary chondrocytes after XJB-5-131 treatment with or without TBHP (n = 5). (B) Flow cytometry analysis and quantitative analysis of intracellular total reactive oxygen species (ROS), FerroOrange (Fe<sup>2+</sup>), lipid peroxidation (C11-BODIPY) (n = 6). (C) Q-PCR analysis of indicated genes in chondrocytes treated with 50  $\mu$ M TBHP and/or 2  $\mu$ M XJB-5-131 for 6 h (n = 5). (D) Western blot analysis of protein expression and quantification of Col2a1, Cox2, MMP13, Fth1 and Gpx4 in chondrocytes treated with XJB-5-131 (2  $\mu$ M) with or without TBHP (n = 3). The values are presented as the means  $\pm$  SDs. \*P < 0.05; \*\*P < 0.01; \*\*\*P < 0.001.

thickness increased significantly in the knee joint of DMM mice with micro-CT both in 4 weeks and 8 weeks group (Fig. 2A–C). In 4 weeks group, we found the bone volume (BV), subchondral bone mineral density (BMD), bone volume/tissue volume (BV/TV), subchondral trabecular thickness (Tb.Th), the number of osteophyte and subchondral bone plate (SBP) thickness were significantly higher in the DMM mice, while the subchondral trabecular separation (Tb.Sp) significantly

reduced and trabecular number (Tb.N) showed no significant change (Fig. 2A). Intra-articular administration of XJB-5-131 significantly alleviate osteophyte formation and subchondral bone thickness (Fig. 2B). The trend of these index in 8 weeks group were similar to that in 4 weeks group and the effect of XJB-5-131 was more pronounced than that in 4 weeks (Fig. 2D). These results suggest that XJB-5-131 suppresses the osteophyte formation and subchondral bone sclerosis in



**Fig. 2.** XJB-5-131 alleviates DMM-induced bone redundancy formation and subchondral bone microarchitectural changes. (A, C) 3D reconstructed images of mice knee joints and the sagittal view of the medial joint compartment revealing the changes to femoral and tibial surfaces and subchondral bone plate (SBP) thickness, respectively. Red line indicates the thickness of SBP (4 weeks and 8 weeks, n = 6). (B, D) Micro-CT analysis of indicated markers of bone redundancy formation and subchondral bone microarchitectural changes (4 weeks and 8 weeks, n = 6). Scale bars, 1 mm (A, C) The values are presented as the means ± SDs. \*P < 0.05; \*\*P < 0.01; \*\*\*P < 0.001.

DMM-induced mice OA model.

### 3.3. XJB-5-131 alleviates OA progression via inhibiting chondrocytes ferroptosis

Next, we performed safranin O/fast green staining of mice joints and found decreased cartilage thickness, reduced chondrocytes numbers, and increased OARSI scores in the DMM mice compared to the Sham mice in both 4 weeks and 8 weeks group, which were significantly alleviated by the intra-articular injection of XJB-5-131 (Fig. 3A–D). To further evaluate the anti-ferroptotic and chondroprotective effects of XJB-5-131, we performed immunofluorescence staining of mice articular cartilage. The histological examination revealed reduced Col2a1, Fth1 and Gpx4 expression, whereas increased Mmp13 and Cox2 expression in DMM-induced mice cartilage, which were markedly abolished by XJB-5-131 treatment both in 4 weeks and 8 weeks group (Fig. 3E–G). Taken together, these data indicates that XJB-5-131 inhibits DMM-induced articular cartilage erosion and chondrocyte ferroptosis.

### 3.4. Pebp1 is a potential mediator of XJB-5-131 exerting the anti-ferroptotic effect in chondrocytes

To explore the mechanisms of XJB-5-131 inhibiting chondrocytes ferroptosis, we performed RNA sequencing analysis of mouse chondrocytes treated with TBHP in the presence or absence of XJB-5-131. Compared with control group, there were 8361 DEGs in TBHP group, within which 4239 genes were up-regulated and 4122 were down-regulated. Moreover, compared with the TBHP group, there were 5500 differentially expressed genes in the TBHP + XJB-5-131 group, of which 2567 were up-regulated and 2933 were down-regulated (Fig. 4A). There were 3870 common difference genes between Control group vs TBHP group and TBHP group vs TBHP + XJB-5-131 group (Fig. 4B). Compared with the Control group, the GO in TBHP group was significantly enriched ROS metabolic processes, cellular transition metal ion homeostasis and regulation of membrane lipid distribution (Fig. 4C). Furthermore, the cellular iron ion homeostasis and glutathione peroxidase activity were more enriched while the ROS metabolic processes were less enriched in the TBHP + XJB group than the TBHP group (Fig. 4C). All of these reveals that XJB-5-131 significantly suppressed TBHP induced chondrocyte ferroptosis.

Within the heat map of DEGs, we noted that the Pebp1 was significantly down-regulated after TBHP treatment and was restored by XJB-5-131 (Fig. 4D). Therefore, we speculated that Pebp1 may be an important mediator of XJB-5-131 resistance to chondrocyte ferroptosis. Then, we performed immunofluorescence staining of mice knee sections to verify the expression of Pebp1 protein in cartilage. Both in 4 weeks and 8 weeks groups, the Pebp1 were significantly decreased in the DMM mice while XJB-5-131 treatment evidently restored its expression (Fig. 4E). Taken together, these results suggest that Pebp1 is a potential target of XJB-5-131 exerting anti-ferroptotic effects in chondrocytes.

### 3.5. Inhibition of pebp1 disables the effect of xjb-5-131 resistant to chondrocyte ferroptosis

Pebp1 was a small hybrid scaffold protein, which was bound to and inhibited the Raf1 kinase cascade under homeostatic conditions, therefore, Pebp1 was also known as Raf1 kinase inhibitory protein (RKIP) [28]. Locostatin, a Pebp1 specific inhibitor, disrupted the interaction of Pebp1 and Raf1 kinase and liberated Pebp1 by binding 15LO [29,30]. We selected Locostatin as an inhibitor of Pebp1 to explore the mechanism of Pebp1 in XJB-5-131 against chondrocyte ferroptosis. We treated primary chondrocytes with different concentrations of Locostatin for 24 h and revealed no significant effect on chondrocyte viability at 5 and 10  $\mu$ M. In contrast, the same concentration of Locostatin blocked XJB-5-131 from rescuing the TBHP induced chondrocyte toxicity (Fig. 5A). Next, we found Locostatin (10  $\mu$ M) attenuates the beneficial effects of

XJB-5-131 on alleviating the ferroptotic hallmarks with flow cytometry (Fig. 5B). Furthermore, Locostatin counteracted XJB-5-131's restoration of cartilage anabolic genes (Col2a1, Acan) and ferroptosis suppressor gene (Gpx4) as well as inhibition of cartilage catabolic genes (Mmp3, Mmp13) and ferroptosis drivers (Atf3, Ptgs2). In addition, the Pebp1 expression was decreased significantly after Locostatin treatment (Fig. 5C). Furthermore, similar results of cartilage metabolic, ferroptosis markers, and Pebp1 were also observed by western blot analysis (Fig. 5D). These results indicates that inhibition of Pebp1 disables the effect of XJB-5-131 resistance to chondrocyte ferroptosis.

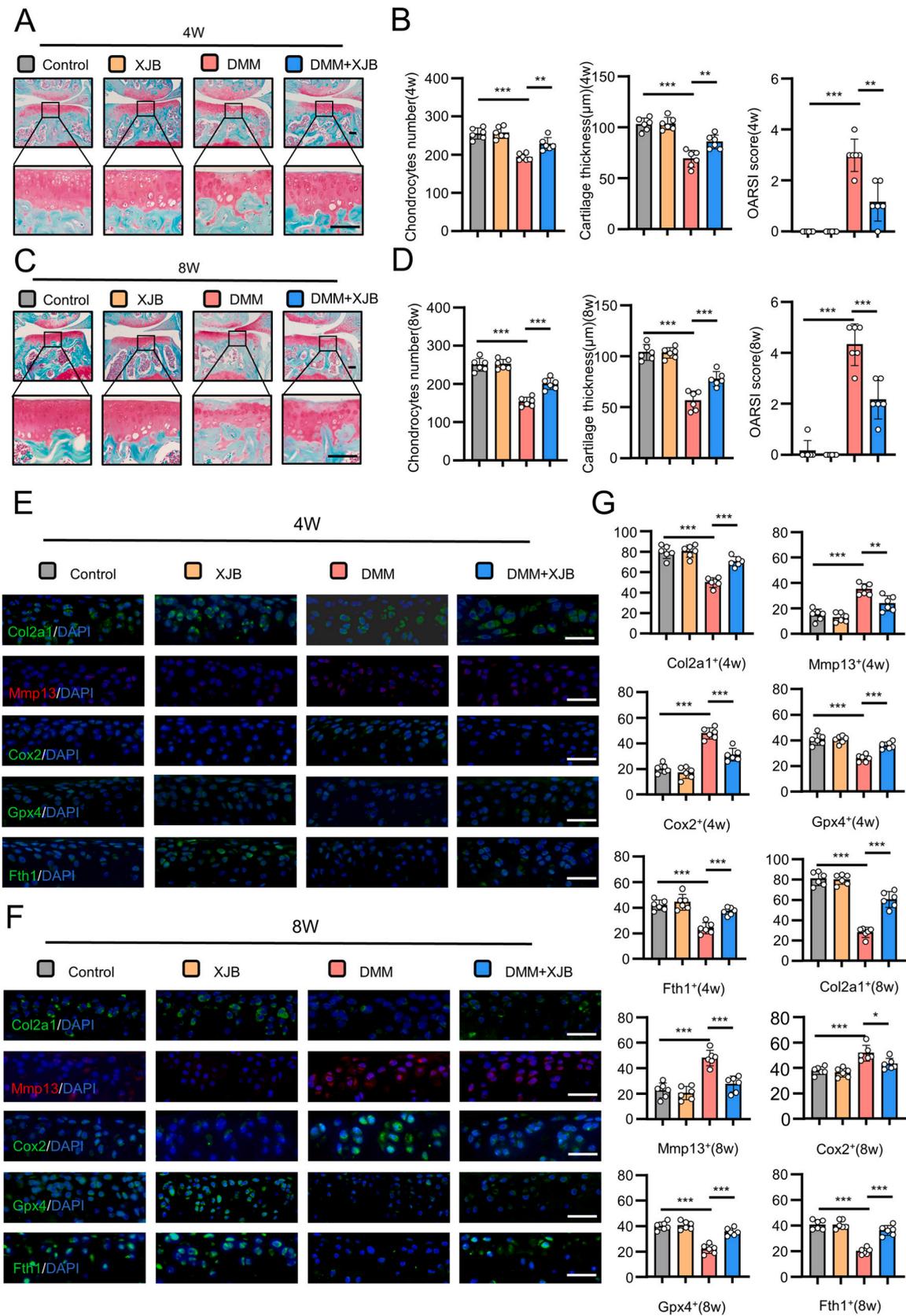
## 4. Discussion

In this study, we explored the effect of XJB-5-131 in inhibiting chondrocyte ferroptosis and alleviating OA progression. There were several novel and important findings here: (1) XJB-5-131 significantly alleviated chondrocyte ferroptosis and restored cartilage metabolic homeostasis in vitro; (2) XJB-5-131 alleviated articular cartilage degeneration in DMM surgery induced OA mice model by suppressing chondrocyte ferroptosis; (3) XJB-5-131 exerted anti-ferroptotic effect on chondrocytes by restoring Pebp1 expression.

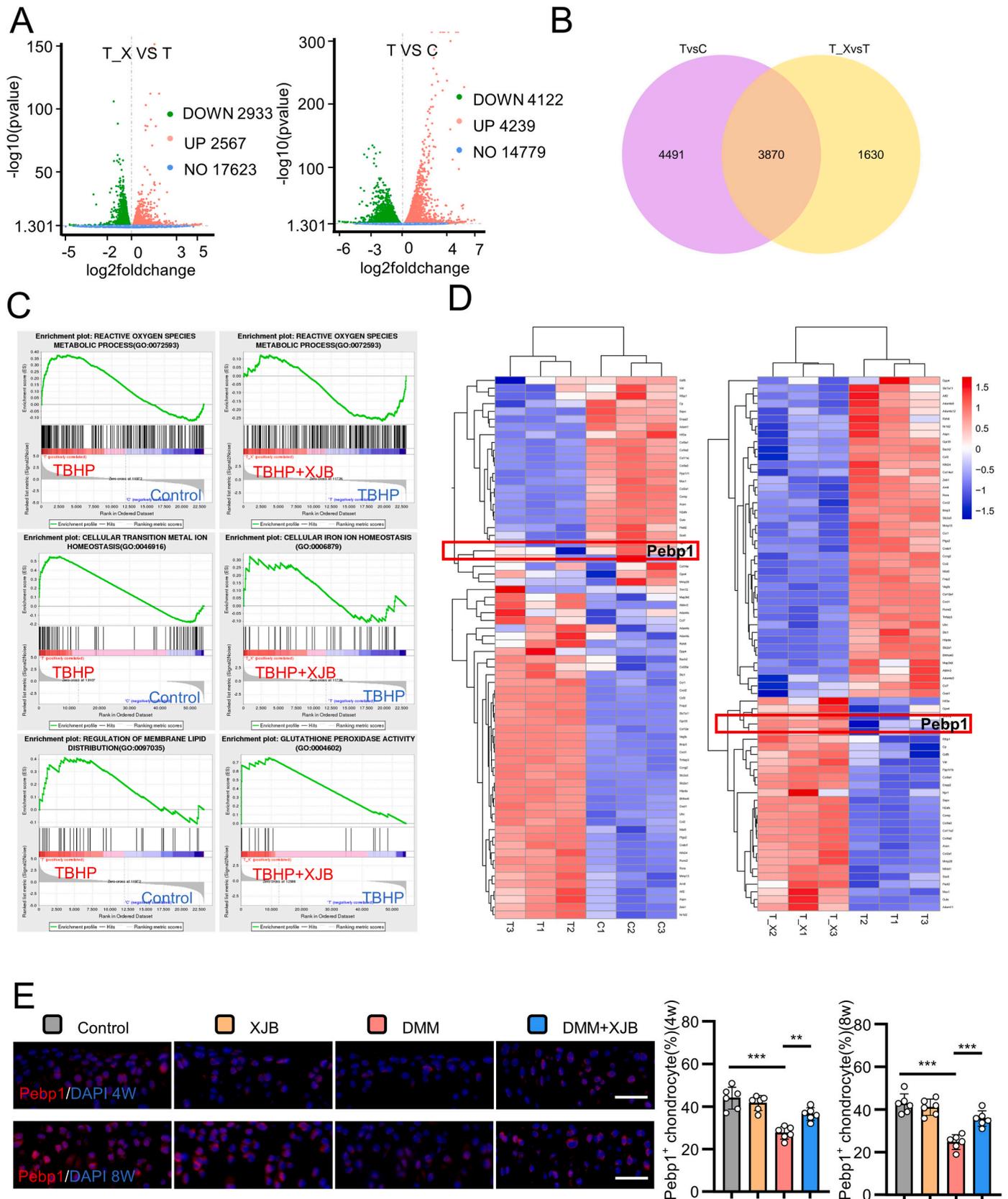
Recent studies have shown that ferroptosis plays an important role in the pathogenesis of various diseases, including OA [31–34]. Iron was not only an essential trace element in human body, but also indispensable for metabolic reactions in chondrocytes, such as redox reactions, cellular respiration and DNA synthesis [35,36]. However, the range of cellular benefits of iron was narrow, and intracellular iron overload in chondrocytes had been suggested as a risk factor for OA [37]. Iron in cells was mainly stored in ferritin, which consists of two types, ferritin heavy polypeptide 1 (Fth1) and ferritin light polypeptide 1 (Ftl1), in which Fth1 played the main role of iron storage [38]. Studies showed that Fth1 reduction would lead to the accumulation of Fe<sup>2+</sup> and subsequent Fenton reactions, which would trigger ferroptosis. Conversely, increased Fth1 inhibited ferroptosis [39–41]. After treatment of XJB-5-131, the expression of Fth1 and Gpx4 increased while the ferroptosis driver Cox2 decreased significantly in primary chondrocytes. In the animal study, we could not obtain chondrocytes and use probes to detect the biochemical markers (ROS levels and lipid peroxidation) of ferroptosis of chondrocytes directly because various factors including temperature and digestive enzymes would interfere the chondrocytes state during the process of cartilage acquisition, resulting in unreliable results. Therefore, immunofluorescence assay was used to measure the markers of cartilage synthesis and catabolism as well as ferroptosis to evaluate the anti-ferroptotic role and cartilage protective effect of XJB-5-131.

We screened and verified Pebp1 as a potential mediator of XJB-5-131 resistance to ferroptosis with RNA sequencing analysis and histological studies. Pebp1 was suggested as a key factor in the regulation of ferroptosis by binding with 15-lipoxygenase (15LOX) to form 15LOX/Pebp1 complex [42–45]. Wenzel et al. suggested that Locostatin disrupted the binding of Pebp1 to Raf1, and liberated it to form 15LOX/Pebp1 complex to exacerbate ferroptosis [42]. In our study, the anti-ferroptotic effect of XJB-5-131 was significantly inhibited by Locostatin. XJB-5-131 protects chondrocytes from ferroptosis by restoring Pebp1 expression, and its inhibition will lead to the failure of XJB-5-131 treatment.

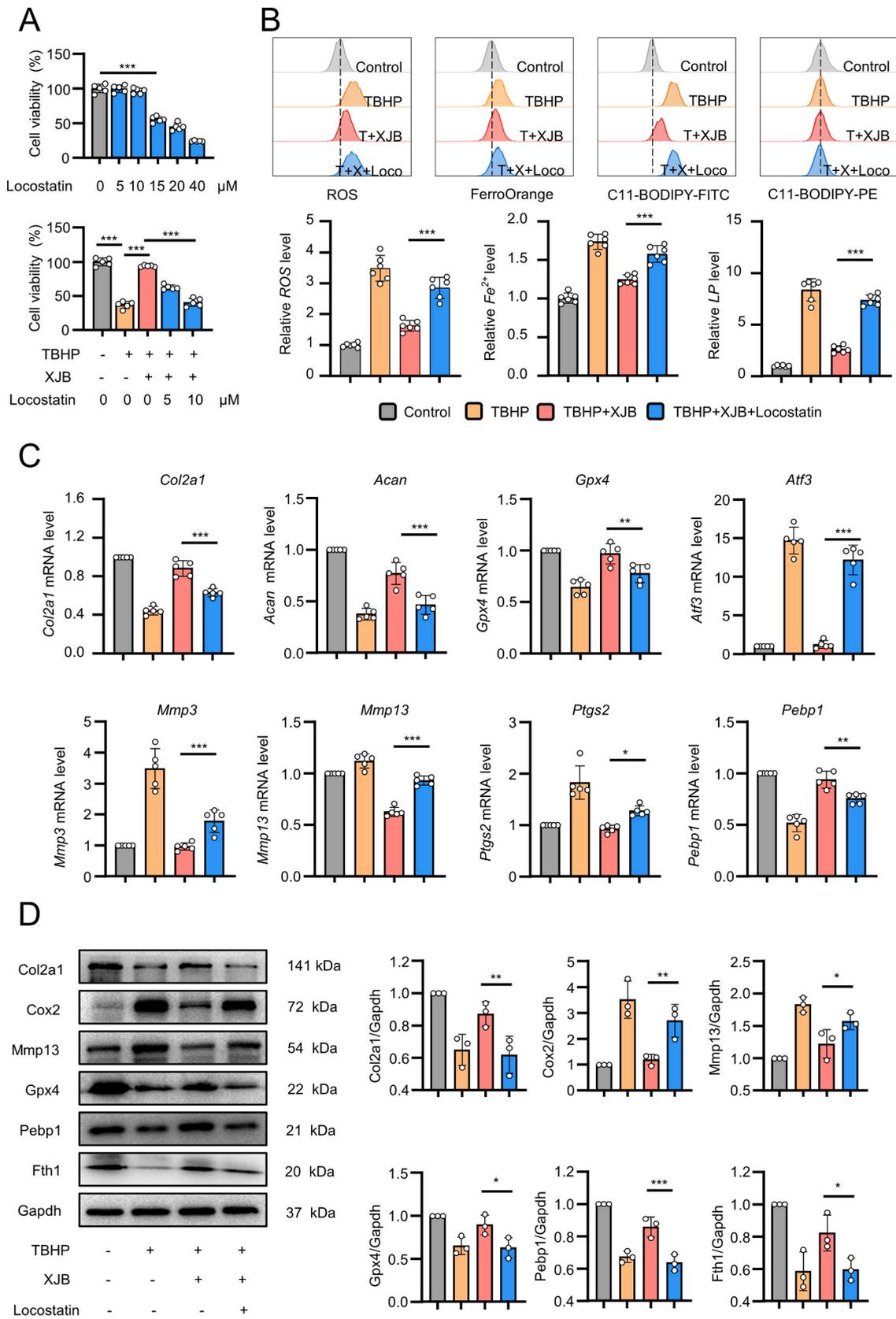
A growing number of studies have targeted the inhibition of chondrocyte ferroptosis since it was found to be involved in the progression of osteoarthritis [10,46,47]. However, there was few studies on targeting mitochondria to inhibit chondrocyte ferroptosis. The function of mitochondria in ferroptosis has been extensively studied, the main opinion was that mitochondria was important source of ROS involved in ferroptosis [48]. Moreover, ROS would lead to mitochondrial membrane lipid peroxidation [49,50]. The mitochondria-targeted antioxidant XJB-5-131 exhibited a potent anti-ferroptotic effect in cardiac cells and tubular epithelial cells [23,51]. Our investigation directly revealed that XJB-5-131 could alleviate OA progression by inhibiting chondrocyte



**Fig. 3.** XJB-5-131 ameliorates DMM-induced cartilage erosion and chondrocyte ferroptosis in mouse knee joints. (A, C) Representative images of safranin-O/fast green stained knee joint sections in different groups (4 weeks and 8 weeks). (B, D) Quantitation of chondrocytes numbers, cartilage thickness and Osteoarthritis Research Society International (OARSI) scores (4 weeks and 8 weeks, n = 6). (E, F) IF staining of Col2a1, Mmp13, Cox2, Fth1, Gpx4 and their quantification (G) in different mice groups (4 weeks and 8 weeks, n = 6). Scale bars, 50 μm (A) (C) (E) (F). The values are presented as the means ± SDs. \*P < 0.05; \*\*P < 0.01; \*\*\*P < 0.001.



**Fig. 4.** RNA sequencing analysis of primary chondrocytes treated with TBHP in the presence or absence of XJB-5-131, which identified Pebp1 as a potential mediator of XJB-5-131 exerting the anti-ferroptosis in chondrocytes. (A) RNA sequencing Volcano Plot between TBHP group (T) vs TBHP + XJB-5-131 group (T\_X) and Control Group (C) vs T. (B) Venn Diagram of C vs T and T vs T\_X (C) Gene ontology (GO) enrichment plot in T vs C and T vs T\_X groups. (D) Heatmap of Differential gene, identified Pebp1 as a potential mediator (E) IF staining of Pebp1 and quantification in different mice groups (4 weeks and 8 weeks, n = 6). Scale bars, 50 μm (E).



**Fig. 5.** Inhibition of pebp1 disables XJB-5-131 resistance to chondrocyte ferroptosis. (A) The viability of primary chondrocytes after Locostatin treatment with or without TBHP and XJB-5-131(n = 5). (B) Flow cytometry analysis and quantitative analysis of intracellular ROS, FerroOrange (Fe<sup>2+</sup>), lipid peroxidation (C11-BODIPY) in different groups(n = 6). (C) Q-PCR analysis of indicated genes in chondrocytes treated with TBHP, TBHP + XJB-5-131 and TBHP + XJB-5-131+Locostatin for 6 h (n = 5). (D)Western blot analysis of protein expression and quantification of Col2a1, Cox2, MMP13, Fth1 and Gpx4 in different groups (n = 3). The values are presented as the means ± SDs. \*P < 0.05; \*\*P < 0.01; \*\*\*P < 0.001.

ferroptosis. We not only provided a potential possibility of modifying the pathological progression of OA but also indicated the direction for the subsequent application of XJB-5-131 as a ferroptosis inhibitor in other diseases. Nonetheless, there were several limitations in our study. OA was a multifactorial degenerative disease involving multiple cells and factors [52,53]. However, in this study, we only investigated the pathogenic mechanism of XJB-5-131 on ferroptosis in chondrocytes, while the studies on other cells (such as macrophage) and other pathogenic factors need to further studied. In addition, the pathological features of OA were degeneration of articular cartilage, hyperplasia of subchondral bone, osteogenesis at the joint edge, inflammation and proliferation of synovial tissue [54]. In our study, although we have provided microCT data of subchondral bone which is consistent with the pathological alleviation of OA, ferroptosis is related to bone homeostasis [55], and the specific mechanism of subchondral bone remodeling needs to be further explored.

## 5. Conclusion

In conclusion, we found that XJB-5-131 suppressed chondrocyte ferroptosis and alleviated OA progression by restoring *Pebp1* expression. This work suggests XJB-5-131 as a potential therapeutic drug for exploring the disease-modifying drug of OA.

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## Declaration of competing interest

A conflict of interest occurs when an individual's objectivity is potentially compromised by a desire for financial gain, prominence, professional advancement or a successful outcome. The Editors of the *Journal of Orthopaedic Translation* strive to ensure that what is published in the Journal is as balanced, objective and evidence-based as possible. Since it can be difficult to distinguish between an actual conflict of interest and a perceived conflict of interest, the Journal requires authors to disclose all and any potential conflicts of interest.

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