Tougu Xiaotong capsule promotes chondrocyte autophagy by regulating the Atg12/LC3 conjugation systems

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Abstract. We have previously reported that Tougu Xiaotong capsule (TXC) inhibits tidemark replication and cartilage degradation by regulating chondrocyte autophagy in vivo. Autophagy, a cell protective mechanism for maintaining cellular homeostasis, has been shown to be a constitutively active and protective process for chondrocyte survival. However, it remains unclear whether TXC promotes chondrocyte autophagy by regulating the autophagy-related (Atg)12/microtubule-associated protein 1 light chain 3 (LC3) conjugation systems. Thus, in the present study, we investigated the effects of TXC on primary chondrocytes treated with cobalt chloride (CoCl2). We found that CoCl2 induced a decrease in chondrocyte viability and the autophagosome formation of chondrocytes, indicating that CoCl2 induced autophagic death in a dose- and time-dependent manner. To determine the effects of TXC on CoCl2-exposed chondrocytes, we assessed cell viability by MTT assay. Our results revealed that TXC enhanced the viability of CoCl2-exposed chondrocytes. To gain insight into the mechanisms responsible for the enhancing effects of TXC on CoCl2-exposed chondrocytes, the expression of Atg genes was assessed in chondrocytes exposed to CoCl2 and treated with or without TXC. The results revealed that the expression of beclin 1, Atg3, Atg5, Atg7, Atg10, Atg12 and LC3 II/LC3 I in the chondrocytes treated with TXC increased, compared to that in the untreated chondrocytes. In addition, ultrastructural analysis indicated that treated chondrocytes contained more autophagosomes than the untreated cells, suggesting that TXC increased the formation of autophagosomes in the chondrocytes to clear the CoCl2-induced autophagic death. Therefore, these data suggest that TXC is a potential therapeutic agent for the reduction of cartilage degradation that occurs in osteoarthritis.

Introduction

Osteoarthritis (OA), a highly prevalent joint disease, exhibits a number of histological characteristics, including a gradual degradation of the extracellular matrix (ECM) and reduced cartilage cellularity, as well as a disruption of the articular cartilage surface, belonging to the GuBi of Traditional Chinese Medicine (TCM) (1-3). Chondrocytes, the only cell population of the articular cartilage, are capable of responding to structural changes in the surrounding ECM by maintaining the dynamic equilibrium between production of the ECM and its enzymatic degradation; however, the capacity of chondrocytes to regenerate the normal ECM architecture is limited and declines with aging due to abnormal responsiveness to anabolic stimuli and cell death (4,5). A number of factors may be involved in the development of OA; however, one of the most important risk factors is cell death. Cell death diminishes the ability of cells to proliferate, mainly due to an increase in apoptosis, which is thought to be a major cause of chondrocyte depletion during OA progression. Several studies have demonstrated that another type of cell death, autophagy, is involved in chondrocyte depletion during OA progression (6-8).

Autophagy, a cellular homeostatic mechanism, plays an important role in nutrient and energy regulation, and in targeting dysfunctional and altered cytosolic macromolecules, membranes and organelles for delivery to lysosomes for recycling and degradation (9-11). At the cellular level, failure of autophagy leads to the increased production of abnormal gene expression, reactive oxygen species, and may cause cell death (12). The consequences of autophagy failure at the tissue and organismal level are abnormal skeletal development, cardiomyopathies, neurodegeneration and premature death (7,13,14). The mammalian target of rapamycin (mTOR), an important suppressor of autophagy, functions upstream of the autophagy-related (Atg) proteins and is crucially regulated by multiple upstream signaling pathways involving adenosine monophosphate (AMP)-activated protein kinase and phosphoinositide 3 (PI3)-kinase/Akt (15,16). In articular cartilage, which is characterized by a very low rate of cell turnover, autophagy appears to be essential to maintain cellular integrity, survival and function (7,8). Previous studies have verified that autophagy is a constitutively active and apparently protective process for the homeostatic state in normal cartilage (17). A
reduced expression of Atg genes has been observed in OA in humans and mice and is accompanied by an increase in chondrocyte apoptosis, indicating a protective and survival-promoting function of autophagy (7,8,17). 

Tougu Xiaotong capsule (TXC; Medical License number: MINZHIZI Z20100006) is composed of a combination of 4 natural products, including Morinda officinalis, Radix Paeoniae Alba, Rhizoma Chuanxiong and Sarcandra glabra. According to the theories of TCM, these natural products mixed together confer the TXC properties of nourishing Shen, filling in the trunk, relaxing musculature and collaterals to limber the joints and strengthen bones and tendons. TXC has been widely used for the therapy of OA in the Second People’s Hospital affiliated to Fujian University of TCM for 2 decades and has been shown to control pain and improve dysfunction in patients with OA (18). We have previously reported that TXC inhibits amidase replication and cartilage degradation by the regulation of chondrocyte autophagy (19). However, the precious molecular and cellular mechanisms responsible for the effects of TXC on the regulation of chondrocyte autophagy remain largely unknown. Thus, the aim of this study was to establish a proof-of-principle that the pharmacological enhancement of autophagy may be an effective therapeutic approach for OA by regulating the Atg12/microtubule-associated protein 1 light chain 3 (LC3) conjugation systems. Our data demonstrate that TXC reduces the severity of chondrocyte damage, at least in part by activating autophagy, suggesting that TXC promotes chondrocyte autophagy, contributing to the regulation of cartilage homeostasis.

Materials and methods

TXC extracts and fingerprint analysis. TXC herbs were dried for 24 h in an air-circulating oven at 50°C and then shredded and crushed to an appropriate particle size in a high-speed rotary cutting mill (ZN-400A; Zhongnan Pharmaceutical Machinery Factory, Changsha, China). According to the proportion of TXC (Morinda officinalis:Radix Paeoniae Alba:Rhizoma Chuanxiong:Sarcandra glabra = 2:2:1:1), 108 g of herbal powder were extracted with 1.5 l distilled water by reflux for 2 h at 50°C in an air-circulating oven at 50°C. Thereafter, the medium was aspirated followed by TXC (0, 50, 100, 200, 300, 400 µg/ml) + 100 µM CoCl₂ for 48 h or with 200 µg/ml of TXC + 100 µM CoCl₂ for 48 h. Following treatment, 20 µl MTT stock solution (5 mg/ml) were added to each well, and the cells were incubated at 37°C for 4 h. Thereafter, the medium was aspirated followed by the addition of 200 µl DMSO, and the cells were shaken for 10 min. The color formed was determined by an ELISA plate reader (EXL 800; BioTek, Winooski, VT, USA).

Reverse transcription-polymerase chain reaction (RT-PCR). Total RNA was extracted from the chondrocytes using TRIzol reagent (Invitrogen, Carlsbad, CA, USA). Reverse transcription was performed using random primers and Superscript™ III (Invitrogen). PCR reactions were conducted on an ABI Prism 7900HT Fast Real-Time PCR System (Applied Biosystems, Foster City, CA, USA). Primers for the amplifications were designed as follows: hypoxia-inducible factor-1α (HIF-1α) forward, 5'-GCA TCT CCA CCT TCT ACC C-3' and reverse, 5'-TTC TGC TCC ATT CCA TCC T3'-3'; 386 bp; beclin 1 forward, 5'-GCT CAG TAC CAG CGA GAA TA-3' and reverse, 5'-GTC AGG GAC TCC AGA TAC GA-3', 350 bp; mTOR forward, 5'-GGA CGG TGT AGA ACT TGG A-3' and reverse, 5'-GAG ATG TCG TCT GGT TGA T-3'; 320 bp; Atg3 forward, 5'-GGA GCC TGT AGA ACT TGG A-3' and reverse, 5'-GAG ATG TCG TCT GGT TGA T-3'; 320 bp; Atg3 forward, 5'-GGA GCC TGT AGA ACT TGG A-3' and reverse, 5'-GAG ATG TCG TCT GGT TGA T-3'; 320 bp; Atg4 forward, 5'-GGA GCC TGT AGA ACT TGG A-3' and reverse, 5'-GAG ATG TCG TCT GGT TGA T-3'; 320 bp; Atg5 forward, 5'-GGA GCC TGT AGA ACT TGG A-3' and reverse, 5'-GAG ATG TCG TCT GGT TGA T-3'; 320 bp; Atg6 forward, 5'-GGA GCC TGT AGA ACT TGG A-3' and reverse, 5'-GAG ATG TCG TCT GGT TGA T-3'; 320 bp; Atg7 forward, 5'-GGA GCC TGT AGA ACT TGG A-3' and reverse, 5'-GAG ATG TCG TCT GGT TGA T-3'; 320 bp; Atg8 forward, 5'-GGA GCC TGT AGA ACT TGG A-3' and reverse, 5'-GAG ATG TCG TCT GGT TGA T-3'; 320 bp.
Western blot analysis. Total protein was extracted from the chondrocytes using RIPA buffer, and protein concentrations were determined using a Bio-Rad protein assay (Bio-Rad, Hercules, CA, USA). Total protein (50 µg) was fractionated by SDS-PAGE, and transferred onto a PVDF membrane (Invitrogen). The PVDF membrane was blocked with 5% non-fat milk and incubated with antibodies to HIF-1α, beclin 1, mTOR, Atg3, Atg7, Atg12, LC3 I/II and β-actin (Santa Cruz Biotechnology, Santa Cruz, CA, USA). The blots were developed using a horseradish peroxidase-conjugated secondary antibody (Beyotime Institute of Biotechnology, Shanghai, China). Immunoreactive proteins were visualized by western blotting chemiluminescence luminol reagent (Santa Cruz Biotechnology). Immunoblotting band gray values were calculated using the Tocan 190 protein assay system (Bio-Rad).

Transmission electron microscopy (TEM). The chondrocytes were fixed in 3% glutaraldehyde and 1.5% paraformaldehyde solution (pH 7.3) at 4°C for 24 h, post-fixed with 1% osmic acid and 1.5% potassium hexacyanoferrate (II) solution (pH 7.3) at 4°C for 2 h, rinsed with water, dehydrated in a graded series of ethanol followed by propylene oxide, kept overnight and embedded in Epon-Araldite resin. Ultrathin sections were obtained using a Leica ultramicrotome and stained with 2% aqueous uranyl acetate, counterstained with 0.3% lead citrate and observed under a transmission electron microscope (H7650; Hitachi High-Technologies Corp., Tokyo, Japan).

Statistical analysis. All data were collected from at least 3 independent experiments. Statistical analysis was performed using SPSS 13.0 software. All data are presented as the means ± standard deviation (SD) and analyzed by the Student’s t-test and ANOVA. Statistical significance was set at P<0.05.

Results

Quality control of TXC. Compared to the spectrogram and chromatographic peak of retention time with the reference substance (Fig. 1A), the composition of TXC was identified (Fig. 1B), and contained paeoniflorin, ferulic acid, isofraxidin and rosmarinic acid.

Identification of chondrocytes. Type II collagen has been identified as the major molecular form of collagen in articular cartilage and is responsible for tensile strength, whereas proteoglycans provide the compressive stiffness necessary for normal articulation and function (5). The second generation of chondrocytes was cultured for 3 days followed by type II collagen immunohistochemical staining. The cytoplasm was stained with brown, representing positive expression in chondrocytes (Fig. 2A and B). The second generation of chondrocytes showed a typical morphology of chondrocytes with a polygonal or spherical shape (Fig. 2C and D). Therefore, we used the second generation of chondrocytes in the subsequent experiments.

TXC increases the cell viability of CoCl₂-exposed chondrocytes. CoCl₂ is commonly used to activate autophagic death by inducing HIF-1α. To establish the cell model of autophagic death, the chondrocytes were treated with various concentrations and for different periods of time with CoCl₂ to determine the effective concentration and treatment duration time by MTT assay. As shown in Fig. 3A, in the cells that were treated with CoCl₂ concentrations of 50 µM (68.43±3.78%), 100 µM (50.38±4.12%), 200 µM (42.84±1.95%), 300 µM (41.13±3.20%) and 400 µM (36.10±4.89%) for 24 h, a dose-dependent decrease in cell viability was observed compared to the untreated cells (100±0.00%) (P<0.01). Cell viability gradually decreased with the increase in the duration of treatment in the chondrocytes treated with 100 µM CoCl₂ for different periods of time (Fig. 3B), suggesting that CoCl₂ inhibited cell viability in a dose- and time-dependent manner due to the CoCl₂-induced autophagic death.

To explore the effects of TXC on CoCl₂-treated chondrocytes, we examined the cell viability of CoCl₂-exposed chondrocytes treated with various concentrations of TXC and for different periods of time by MTT assay. As shown in Fig. 3C, the cell viability of the CoCl₂-exposed chondrocytes treated with TXC concentrations of 50 µg/ml (105.65±3.38%), 100 µg/ml (111.38±3.04%), 200 µg/ml (121.43±5.67%), 300 µg/ml (118.44±3.48%) and 400 µg/ml (119.81±2.77%) for 48 h was enhanced in a dose-dependent manner compared to that of the cells treated with CoCl₂ only (100±0.00%) (P<0.01). Cell
viability gradually increased with the increase in the duration of treatment in the CoCl$_2$-exposed chondrocytes treated with 200 µg/ml TXC for different periods of time (Fig. 3D), indicating that TXC promoted the survival of CoCl$_2$-exposed chondrocytes.

**TXC enhances chondrocyte survival by promoting cell autophagy.** TXC has been reported to delay cartilage degradation by activating chondrocyte autophagy in vivo; however, it remains to be seen whether TXC enhances chondrocyte survival by increasing the expression of Atg genes. To verify the effects of TXC on chondrocyte autophagy, we observed the morphology of CoCl$_2$-exposed chondrocytes treated with or without TXC under a phase-contrast microscope.

To explore the role of TXC in chondrocyte autophagy, the mRNA and protein expression of Atg genes in CoCl$_2$-exposed chondrocytes treated with or without TXC was examined by...
**RT-PCR and western blot analysis, respectively.** The results revealed that the mRNA expression of beclin 1, Atg3, Atg5, Atg7, Atg10, Atg12 and LC3 II/LC3 I in the chondrocytes treated with TXC increased, compared to the untreated cells (P<0.05, P<0.01) (Fig. 5). The protein expression levels of these Atg genes were similar to their respective mRNA expression levels (Fig. 6), suggesting that TXC promoted chondrocyte autophagy by regulating the Atg12/LC3 conjugation systems.

**Discussion**

The present study systematically investigated the effects of TXC on CoCl2-induced chondrocyte autophagic death in vitro. Our results clearly demonstrated that TXC enhanced the viability of chondrocytes exposed to CoCl2 and increased the formation of autophagosomes in the CoCl2-exposed chondrocytes by upregulating the expression of beclin 1, Atg3, Atg5, Atg7, Atg10, Atg12 and LC3 II/LC3 I. Taken together, these results indicate that TXC is a potential therapeutic agent for the reduction of cartilage degradation that occurs in OA.

Current treatments for the management of OA do not reverse the degradation process of articular cartilage. Non-steroidal anti-inflammatory drugs (NSAIDs) have been used in the treatment of OA for the past several years; however, their therapeutic effects remain unsatisfactory due to the serious adverse side-effects, such as gastrointestinal and cardiovascular diseases (21-23). Accordingly, the development of novel drugs from natural herbs, which can provide cartilage protection and be safely used in the prolonged treatment of OA, is required. TXC has been used to control tidemark replication and cartilage degradation by inhibiting chondrocyte apoptosis and promoting chondrocyte autophagy (18,19). Therefore, in the present study, we sought to determine the efficacy of TXC on chondrocyte autophagy, as a chondroprotective agent.
Excessive mechanical loading of articular cartilage causes damage to chondrocytes and the ECM, and initiates a pathological process that occurs due to abnormal chondrocyte activation and results in the spreading of chondrocyte death and damage to the ECM beyond the initial area that was exposed to the highest mechanical load (24, 25). The understanding of the mechanisms involved in these cellular changes may provide the potential to identify targets for pharmacological interventions in order to attenuate or prevent the subsequent development of OA. In the articular cartilage, autophagy activated by different types of stress has been shown to be a constitutively active and protective process for the survival of chondrocytes (26). A previous study reported that the expression of 3 markers for different stages of autophagy, including uncoordinated-51-like kinase (ULK), beclin 1 and LC3, was decreased in the cartilage of patients with OA and in a mouse model of OA (8).

In the present study, we used a CoCl₂-induced model of cell autophagic death to assess the molecular mechanisms responsible for the promoting effects of TXC on chondrocyte autophagy in vitro. In order to determine the inducer concentration of CoCl₂ in chondrocytes, cell viability was examined. Following exposure to CoCl₂ at various concentrations and for different periods of time, cell viability was inhibited due to the CoCl₂-induced autophagic death as shown by MTT assay. Our results revealed that CoCl₂ induced autophagic death in a dose- and time-dependent manner. The changes observed in cell morphology suggested that the cells underwent autophagic death 24 h following incubation with...
the concentration of CoCl$_2$ selected based on the results of MTT assay. To determine the effective concentration of TXC on the viability of CoCl$_2$-exposed chondrocytes, MTT assay was performed. These results indicated that TXC enhanced chondrocyte survival, as shown by the increased cell viability.

Autophagy, which is in part related to the reduced expression of autophagic regulators, is compromised in osteoarthritic cartilage. In articular cartilage, autophagy does not only occur in response to mechanical injury, but is also deficient with aging (8,27). Autophagy is characterized by the formation of autophagosomes and their fusion with lysosomes. At the late stage of autophagy, lysosomes fuse with and release lysosomal enzymes into the autophagosome to degrade its contents (28). Atg genes control the autophagy process leading to the induction and nucleation of autophagic vesicles, their fusion and expansion with lysosomes, following enzymatic degradation and recycling (29,30). Atg12 undergoes a ubiquitin-like conjugation to Atg5 through an internal lysine residue and a COOH-terminal glycine, respectively. This process is activated by the Atg7 protein, which is homologous to the E1 family of ubiquitin-activating enzymes, and Atg10, which functions as a protein-conjugating enzyme (31). The Atg12-Atg5 conjugates recruit Atg16 dimers. Atg16 is a bivalent molecule, which leads to the formation of large multimeric complexes, and these are thought to play a key role in the nucleation of both cytoplasm-to-vacuole targeting vesicles and autophagosomes (32). The number of lysosomes increases in the process of autophagy, accompanied by an increased expression of beclin 1 and LC3 (33). Beclin 1, forming a complex with type III PI3 kinase and Vps34, participates in the nucleation of the autophagic vesicle (19). The involvement of LC3 in the protein conjugation system is required in the expansion of the autophagosome. There are 2 forms of LC3, including LC3 I, in the cytoplasm and LC3 II bound to the autophagosome membrane. During autophagy, LC3 I is converted to LC3 II by the ubiquitin-like system. Thus, the level of LC3 II and the ratio of LC3 II to LC3 I closely reflect the extent of autophagy (34).

To gain insight into the mechanisms responsible for the effects of TXC on CoCl$_2$-induced autophagic death, the expression of mTOR and HIF-1α, as modulators of autophagy, was assessed in chondrocytes. mTOR plays an important role in multiple cellular functions, such as cell metabolism, proliferation and autophagy (35). All Atg genes have been shown to act downstream of mTOR. HIF-1α, a heterodimeric transcription factor that mediates adaptive responses to hypoxia, serves to regulate both autophagy and apoptosis (36). The present results indicated that the expression of beclin 1, Atg3, Atg5, Atg7, Atg10, Atg12 and LC3 II/LC3 I in the CoCl$_2$-exposed chondrocytes treated with or without TXC significantly increased, compared to the untreated cells. In addition, ultrastructural analysis revealed that the chondrocytes treated with TXC contained more autophagosomes than the untreated cells, suggesting that TXC increases the formation of autophagosomes in chondrocytes to clear the CoCl$_2$-induced autophagic death.

Based on these results, we hypothesized that TXC promotes chondrocyte autophagy by regulating the Atg12/LC3 conjugation systems. Since autophagy serves to delay the onset of apoptosis, experiments are currently in progress in order to explore whether there is a direct association between the induction of autophagy and apoptosis.

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