



# Urotensin-II Prevents Cartilage Degeneration in a Monosodium Iodoacetate-Induced Rat Model of Osteoarthritis

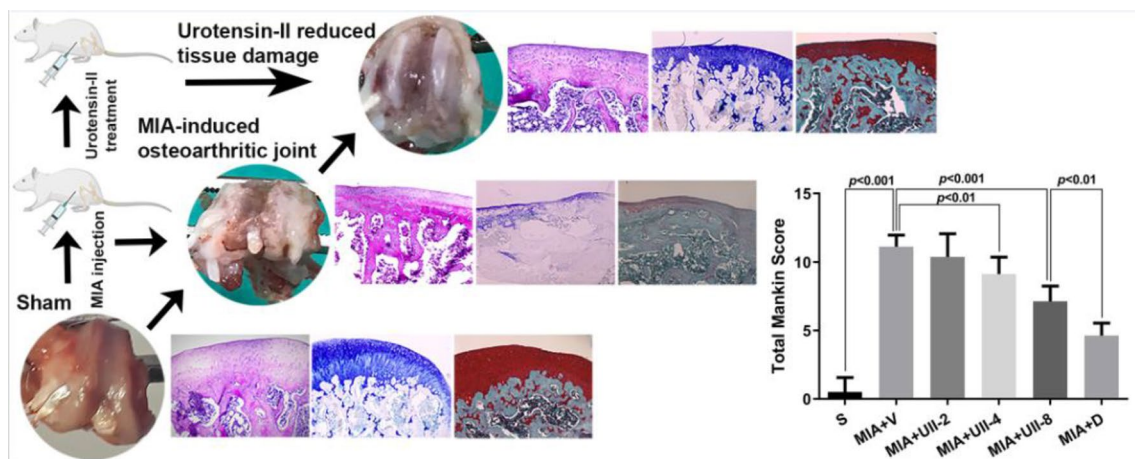
Menderes Yusuf Terzi<sup>1,2</sup> · Hamza Malik Okuyan<sup>3</sup> · İhsan Karaboğaç<sup>4</sup> · Cemil Emre Gökdemir<sup>5</sup> · Duygu Tap<sup>2</sup> · Aydiner Kalacı<sup>5</sup>

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## Abstract

Osteoarthritis (OA) is a common degenerative articular disorder caused by traumatic or spontaneous factors such as genetics, obesity, and advanced age. Comprehending the pathogenic mechanism of OA ensures the development of novel disease-modifying therapeutics rather than conventional palliative drugs with undesired side effects. Urotensin-II (UII) is a multi-functional short cyclic peptide implicated in several disorders. We aimed to analyze the effects of intraarticular UII treatment in a monosodium iodoacetate (MIA)-induced OA rat model. We divided animals into six groups to test three different concentrations of UII with histopathological and immunohistochemical analyses of bone morphogenetic protein-2 (BMP-2), nuclear factor kappa B subunit 1 (NF-κB), and intrinsic UII expression. We analyzed serum levels of cartilage related and inflammatory markers post-OA. We observed a noticeable amelioration of the MIA-induced knee damage in UII-treated animals after gross morphology examination. Mankin scoring after histopathological stainings revealed a partial prevention of articular tissue damage in UII-treated animals. We found a significant reduction in BMP-2 and NF-κB while an increase in intrinsic UII expressions upon exogenous UII injection after immunohistochemical analyses. The Mankin scores were significantly correlated with BMP-2, NF-κB, and intrinsic UII levels. There was no significant alteration in serum markers after UII treatment. We are the first group showing the protective effect of UII on the destructed knee joints of osteoarthritic rats by downregulating the BMP-2 and NF-κB and upregulating intrinsic UII expressions. To uncover the mechanistic role of UII during OA, further experiments are warranted.

## Graphical Abstract



**Keywords** Osteoarthritis · Urotensin-II · Cartilage destruction · Monosodium iodoacetate · BMP-2 · NF-κB

Extended author information available on the last page of the article

## Abbreviations

OA	Osteoarthritis
UII	Urotensin-II
BMP-2	Bone morphogenetic protein 2
NF- $\kappa$ B	Nuclear factor kappa B subunit 1
COMP	Cartilage oligomeric matrix protein
IL-6	Interleukin-6
IL-1 $\beta$	Interleukin-1 $\beta$
ELISA	Enzyme-linked immunosorbent assay

## Introduction

Osteoarthritis (OA) is a prevalent arthritic disorder affecting more than 300 million people worldwide annually (Disease et al. 2018). The incidence of aging-related disorders including OA is projected to rise in parallel with prolonged life expectancy owing to the improvements in the healthcare system (Lunenfeld and Stratton 2013). The main pathological features of OA are cartilage destruction, synovial inflammation, fibrillation, ulceration, and degradation of the articular matrix, and destruction of the joint upper layer, which results in osteophyte formation, the thickness of subchondral bone, and the resultant pain (Felson 2006; Mobasher and Batt 2016). Although several risk factors including genetics, aging, obesity, and trauma have been known to contribute to the onset of OA, its molecular pathogenesis has not been fully understood yet (Geyer and Schonfeld 2018).

OA animal models are common tools to study the pathophysiology of the disease that is a surrogate of human OA. Monosodium iodoacetate (MIA) functions as glyceraldehyde-3-phosphate dehydrogenase inhibitor that hinders glycolysis and causes cell death in chondrocytes (Kuyinu et al. 2016). MIA-induced OA model belongs to the chemically induced OA and is commonly preferred by researchers over other models (spontaneous or traumatic) due to its reproducibility (Teeple et al. 2013).

Today, the therapeutic options for the treatment of OA are aiming at either relieving the pain with drugs as a palliative option or lasting operational joint replacement due to the lack of non-invasive options for permanent rehabilitation (Cao et al. 2020). Therefore, understanding the molecular pathogenesis underlying OA and discovering the disease-modifying therapies are essential to improve the quality of life of the patients with OA and to reduce the socioeconomic costs of OA (Kim et al. 2018).

Urotensin-II (UII) is an 11–17 amino acid-long peptide with a functional cyclic part that is conserved among various species (Sun and Liu 2019). UII plays a vital role in the vascular system by functioning as a strong vasoconstrictor or vasodilator, which is abundantly expressed in the cardiovascular system, kidneys, and central nervous system amongst other tissues and organs (Ross et al. 2010). UII is involved

in not only vaso-modulatory activities but also several physiological and pathological conditions (Sun and Liu 2019). UII and its G protein-coupled receptor-14 (UTR) have been implicated in the pathways pertaining to the inflammatory responses of atherosclerosis and colonic inflammation through bone morphogenetic protein-2 (BMP-2) and nuclear factor kappa B subunit 1 (NF- $\kappa$ B) pathways (McDonald et al. 2007; Yang et al. 2016). Besides, the increased levels of UII in synovial fluids (SF) of OA patients were also reported in a previous clinical study (Gogebakan et al. 2014). Previous studies investigating the action mechanisms of UII highlight the diagnostic and therapeutic importance of this molecule in pathophysiological conditions (Ross et al. 2010; Pereira-Castro et al. 2019). Of which, some studies claim that UII contribute and/or exacerbate fibrosis via mostly recruiting or corroborating the inflammatory pathways in different disease models affecting lung, liver, and heart (Zhang et al. 2007; Kemp et al. 2009; Onat et al. 2012). On the contrary, some other studies report that UII exhibits protective effects mainly by attenuating apoptosis in kidney, heart, and endothelial cells (Chen et al. 2012, 2014; Hsu et al. 2013). However, the underlying action mechanism of this multifunctional vasoactive peptide is still obscure during OA pathogenesis. In this present study, we aimed to investigate the potential therapeutic effects of exogenous UII administration articularly in MIA-induced OA rat model. For this objective, we conducted analyses to check histopathological and immunohistochemical alterations in the central molecules pertaining to inflammation and chondrocyte and bone metabolisms pathways i.e., BMP-2 and NF- $\kappa$ B (Goldring and Marcu 2009) as well as the vasomodulatory peptide UII. We also performed ELISA assays to detect serum levels of prominent inflammatory cytokines interleukin-6 (IL-6) and interleukin-1 $\beta$  (IL-1 $\beta$ ) and a prominent bone/cartilage pathophysiology biomarker; cartilage oligomeric matrix protein (COMP) after intra-articular UII treatment.

## Methodology

### Experimental Animals

All experimental procedures in the present work were abided by the “Animal Care and Use Committee Guidelines” of Hatay Mustafa Kemal University as an act of animal protection utilized for scientific aims. Before the experiments, institutional ethical approval was acquired from the “Animal Ethics Committee of Hatay Mustafa Kemal University” (2020/4–18). We conducted all experiments with a total of 40 male Wistar albino rats weighing 350–400 g and aging 4 months, which were allowed to reach food and water ad libitum. The animals were kept in solid-bottom cages under 23  $\pm$  2 °C, 60–80% humidity and 12-h light/dark cycle

conditions with regularly refreshed ventilation. Prior to the start of the experiments, all animals were kept in their cages for about one week for environmental adaptation.

### OA Model Induction via MIA Injection and Treatment Regimes

We divided 40 rats into 5 individual cages comprising eight animals per group. In order to model OA in animals, at day 0 (zero), a single dose of 0.1 mg/ $\mu$ l (25  $\mu$ l) MIA (M, Sigma–Aldrich, USA) was administered into the infrapatellar ligament of the right knee with a 26.5-G needle (Guzman et al. 2003) under isoflurane-mediated (Hana Pharm, Korea) anesthesia via inhalation (Bove et al. 2003). Fifteen days after the initial MIA injection (or saline as a vehicle) when the fibrosis starts to progress in the knee joint (Guzman et al. 2003), the animals were treated with UII (U7257, Sigma Aldrich, MO, USA) or saline via intra-articular injection at every 3 days for a total of seven injections (Kim et al. 2016) or with 2 mg/kg diclofenac (D, positive control) subcutaneously on daily basis for 21 days. The animal groups were designated as follow (Fig. 1): Group-1 (S): Sham; a single injection of 25  $\mu$ l saline solution intra-articularly into the right knee and 7 intra-articular injections of 25  $\mu$ l saline; Group-2 (MIA + V): MIA + Vehicle; a single dose of intra-articular MIA injection and 7 intra-articular injections of 25  $\mu$ l saline; Group-3 (MIA + UII-2), Group-4 (MIA + UII-4), and Group-5 (MIA + UII-8): A single dose of MIA via intra-articular injections and 7 injections of 2, 4, and 8  $\mu$ g UII in 25  $\mu$ l saline solution respectively (Wu et al. 2010; Okuyan et al. 2021); Group-6 (MIA + D): MIA + diclofenac; a single dose of MIA via intra-articular injection and daily diclofenac treatment for 21 days.

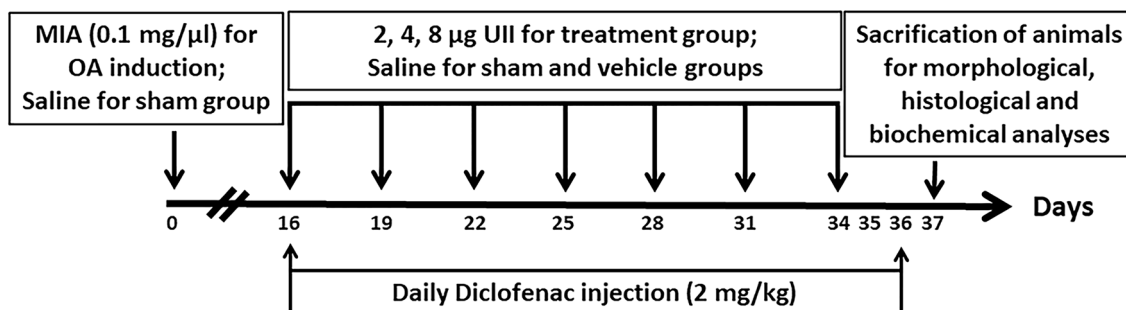
All animals were sacrificed at day 37 via deep anesthesia for further histopathological, immunohistochemical, and ELISA analyses. For the latter, the sera were collected from the whole bloods of the rats via cardiac puncture by spinning down at 2000  $\times$  g for 10 min at 4  $^{\circ}$ C.

### Histopathological Analysis

After the morphological examination, knee samples were put into 10% formalin solution for fixation about 3 days and incubated in 10% EDTA (Sigma Aldrich) for 4 weeks at room temperature for decalcification (Lu et al. 2018). Then, the samples were run through within ascending ethanol series and in absolute xylene for dehydration and finally were embedded in paraffin wax. For histopathological and immunohistochemical analyses, 5- $\mu$ m-thick paraffin sections were sliced with a rotary microtome (CUT-5062, Slee MeDal, Mainz, Germany). Then, the tissue sections were stained with hematoxylin and eosin (H&E) for general cellular structure, safranin O and fast green FCF for collagen and proteoglycan amount, and toluidine blue for glycosaminoglycan amount (Dumond et al. 2004; Lu et al. 2018). Finally, the stained tissue sections were photographed with a light microscope (Olympus CX41, Tokyo, Japan) for further analyses using an image analysis software (Kameram Gen III, Argenit, Istanbul, Turkey). In safranin O & fast green FCF stained slides, medial tibial plateaus were examined for the cartilage degeneration and evaluated using the Mankin scoring system (Mankin 1971) with a scale ranging between 0 and 14 points (0 = No cartilage degeneration, 14 = Severe cartilage degeneration) by two experts blinded to groups (Table 1).

### Immunohistochemical Examination

For NF- $\kappa$ B, BMP-2, and UII immunohistochemical examinations, tissue sections were deparaffinized in xylene, passed through descending ethanol series for rehydration, and rinsed with phosphate-buffered saline (PBS). Then, the sections were washed in 3% hydrogen peroxide solution to block endogenous hydrogen peroxidase activity causing non-specific background staining. Thereafter, the sections were boiled in citrate buffer solution for antigen retrieval. Next, after the slides were washed in PBS and incubated with blocking serum at room temperature for 1 h, they were



**Fig. 1** Design of the experimental procedure. MIA Monosodium iodoacetate, OA Osteoarthritis, UII Urotensin-II

**Table 1** Mankin score grading system for osteoarthritic articular cartilage (Mankin 1971)

Category	Subcategory	Score
Structure	Normal	0
	Surface irregularities	1
	Pannus and surface irregularities	2
	Clefts to transitional zone	3
	Clefts to radial zone	4
	Clefts to calcified zone	5
	Complete disorganisation	6
Cells	Normal	0
	Diffuse hypercellularity	1
	Cloning	2
	Hypocellularity	3
Safranin O staining	Normal	0
	Slight reduction	1
	Moderate reduction	2
	Severe reduction	3
	No dye noted	4
Tidemark integrity	Intact	0
	Crossed by blood vessels	1
Total		0–14

further incubated with anti-BMP-2 (YID0638, YL Biont, Shanghai, China), anti-NF- $\kappa$ B (NB100-56,055, Novus Biologicals, Littleton, CO, USA), and anti-UII (Biorbyt Ltd, Cambridge, UK) primer antibodies (Nf- $\kappa$ B and BMP-2; 1:100, UII; 1:20 dilutions) in a humidified chamber for 60 min at 37 °C. Following incubation, biotinylated secondary antibody and streptavidin peroxidase (Ultra Vision Detection System-HRP kit, Thermo Scientific/Lab Vision, Fremont, CA, USA) were applied to the slides respectively for 20 min at room temperature. Slides were washed in PBS and treated with 3-amino-9-ethylcarbazole (AEC, Thermo Scientific/Lab Vision) as chromogenic substrate. Contrast staining was performed with Mayer's hematoxylin. The UII, NF- $\kappa$ B and BMP-2 stainings were evaluated with an image analysis software (Kameram Gen III, Argenit, Istanbul, Turkey). The staining intensities of UII, NF- $\kappa$ B and BMP-2 were quantitatively assessed by integrated optical density measurement method (Okuyan et al. 2020).

### Measurement of Serum Protein Levels via ELISA

Serum levels of COMP, IL-6, and IL-1 $\beta$  were measured with commercial enzyme-linked immunosorbent assay (ELISA) kits following manufacturers' protocols. The intra-assay and inter-assay coefficients of variations for COMP (Elabscience), IL-6 (Invitrogen), and IL-1 $\beta$  (Invitrogen) were 8% and 5%, 10% and 12%, and 10% and 10% respectively. The sera samples with hemolysis were excluded from the analyses.

### Statistical Analysis

We utilized Shapiro–Wilk test to check whether our data were normally distributed. The significant differences among the groups were tested with parametric one-way ANOVA or non-parametric Kruskal–Wallis H tests. Individual groups were compared via two-way comparison by using non-parametric Mann–Whitney U test or parametric Holm–Sidak's multiple comparison tests. The correlation analyses were conducted with Spearman's correlation test since normal distribution was not met. We utilized SPSS package software (version 22.0) for all statistical analyses and Graphpad (version 8.0.2) for setting plots. The data were expressed as mean  $\pm$  standard deviation or mean  $\pm$  standard error of the mean. We accepted *p* values less than 0.05 as statistically significant.

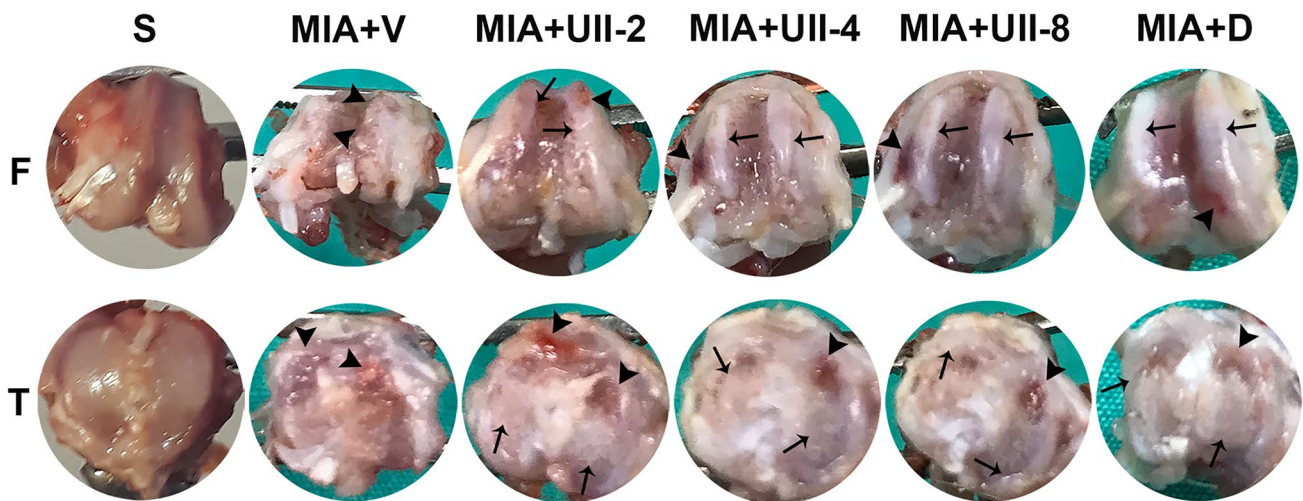
## Results

### Macroscopic Observations

The gross morphologic analyses of the articular surfaces of rats' knee joints were depicted in Fig. 2. MIA administration clearly destructed the cartilage and subchondral bone in MIA + V group compared to the intact and healthy tibial plateau and the femoral condyle in the sham group. UII-treated osteoarthritic rats (MIA + UII-2, -4, and -8) prevented the MIA-induced tissue destruction with a dose-dependent fashion in both cartilage and subchondral bone tissues. The articular knee joints of rats in MIA + D, as the positive control group, also slowed down the tissue destruction inflicted by MIA injection.

### Histopathological Evaluation

In histopathological assessment with H&E staining, the tibial plateau of cartilage surface in the sham group was observed as intact and smooth with a well-stained extracellular matrix (Fig. 3A). Histopathological degeneration was observed in MIA + V group such as rough and irregular cartilage surface, decreased staining density and reduced number of chondrocytes owing to severe osteonecrosis (Fig. 3A). UII treatment prevented the histopathological decay in articular cartilage (Fig. 3A). The toluidine blue and safranin O and fast green FCF stainings clearly expressed an improvement regarding the concentration of glycosaminoglycan (Fig. 3B) and proteoglycan (Fig. 3B, C) in the UII-treated groups compared to the MIA + V group. In OA models, the Mankin scoring system was often used in scoring and classification of histopathological lesions based on staining, structural, and cellular alterations in the articular cartilage (Table 1) (Mankin 1971). Total Mankin



**Fig. 2** Gross morphology of knee joint surfaces; femoral condyles (F) and tibial plateaus (T). Arrowheads indicate the MIA-induced degenerations in cartilage and subchondral bone tissues, whereas arrows

point the improved tissue parts after UII or D treatments. S Sham, MIA Monosodium iodoacetate, V Vehicle, UII-2,4,8 Urotensin-II-2,4,8  $\mu\text{g}$ , D Diclofenac

score and the scores at structure, staining, and cells levels increased in the MIA + V group compared to the sham group indicating the substantial cartilage destruction after MIA injection (Fig. 3D–G,  $p < 0.05$ ). UII administrations led to a significant reduction in total Mankin scores of MIA + UII-4 and MIA + UII-8 groups compared to the MIA + V group (Fig. 3A,  $p < 0.05$ ). Regarding individual scores, while UII-4 and UII-8 significantly reduced the structure and cells scores (Fig. 3F, G,  $p < 0.05$ ), just UII-8 led to significant reduction in staining score compared to the MIA + V group (Fig. 3E,  $p < 0.05$ ). All scores decreased in MIA + D positive control compared to MIA + V group (Fig. 3D–G,  $p < 0.05$ ).

### Immunohistochemical Findings

The effect of UII treatment on the immunohistochemical expressions of BMP-2, NF- $\kappa$ B, and UII was depicted in Fig. 4. The increased immunohistochemical NF- $\kappa$ B expression was observed in MIA + V group compared to the sham group (Fig. 4A, D,  $p < 0.05$ ). Treatment of UII led to a significant decrease in NF- $\kappa$ B expression in MIA + UII-4 and MIA + UII-8 groups compared to MIA + V group (Fig. 4A, D,  $p < 0.05$ ).

The optical intensity of BMP-2 significantly increased in MIA + V group compared to the sham group (Fig. 4B, E,  $p < 0.05$ ). The expression of BMP-2 significantly decreased in MIA + UII-2, MIA + UII-4, and MIA + UII-8 groups compared to the MIA + V group (Fig. 4B, E,  $p < 0.05$ ). Diclofenac treatment significantly reduced both NF- $\kappa$ B and BMP-2 immunoreactivities compared to MIA + UII-8 group (Fig. 4,  $p < 0.05$ ).

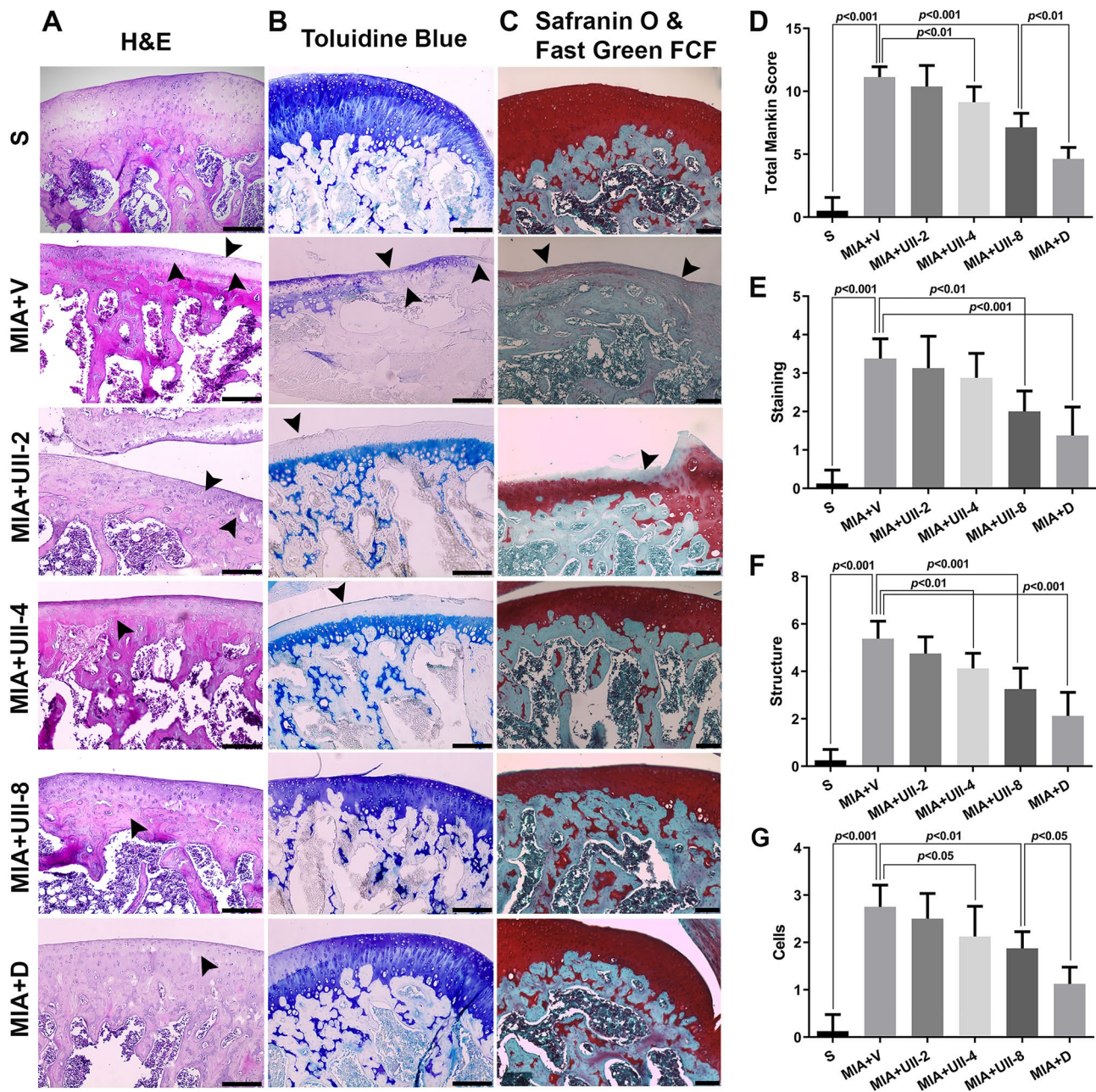
We also performed UII immunohistochemical staining to find out the expression status of UII within articular tissue. We showed that UII is naturally expressed in healthy articular tissue of rats (Fig. 4C). Intra-articular UII expression significantly decreased in MIA + V group compared to the sham group (Fig. 4C, F,  $p < 0.05$ ). We also demonstrated the dose-dependent elevation of intrinsic UII expression after exogenous UII and diclofenac treatments compared to the MIA + V group (Fig. 4C, F,  $p < 0.05$ ).

### Correlation Between Histological Lesion Rate and Tissue-Specific BMP-2, NF- $\kappa$ B, and UII Expressions

BMP-2 is known to be upregulated in osteoarthritic cartilage and responsible for aggrecan degradation and OA progression through close connection with other destructive enzymes (van der Kraan et al. 2010; Liu et al. 2015a, b). In order to analyze any association between the histological recovery rates and the expression levels of BMP-2, NF- $\kappa$ B, and UII in osteoarthritic cartilage after UII treatment, we performed Spearman's correlation test. We found significant correlations between the total Mankin scores and the staining intensities of NF- $\kappa$ B ( $r = 0.8847$ ,  $p < 0.0001$ ), BMP-2 ( $r = 0.9013$ ,  $p < 0.0001$ ), and UII ( $r = -0.9153$ ,  $p < 0.0001$ ) expressions in articular tissues of the rats (Fig. 5).

### Steady Levels of Serum Markers After MIA and UII Treatments

We analyzed the OA-related inflammatory cytokines IL-1 $\beta$  and IL-6 levels together with COMP, as a prominent

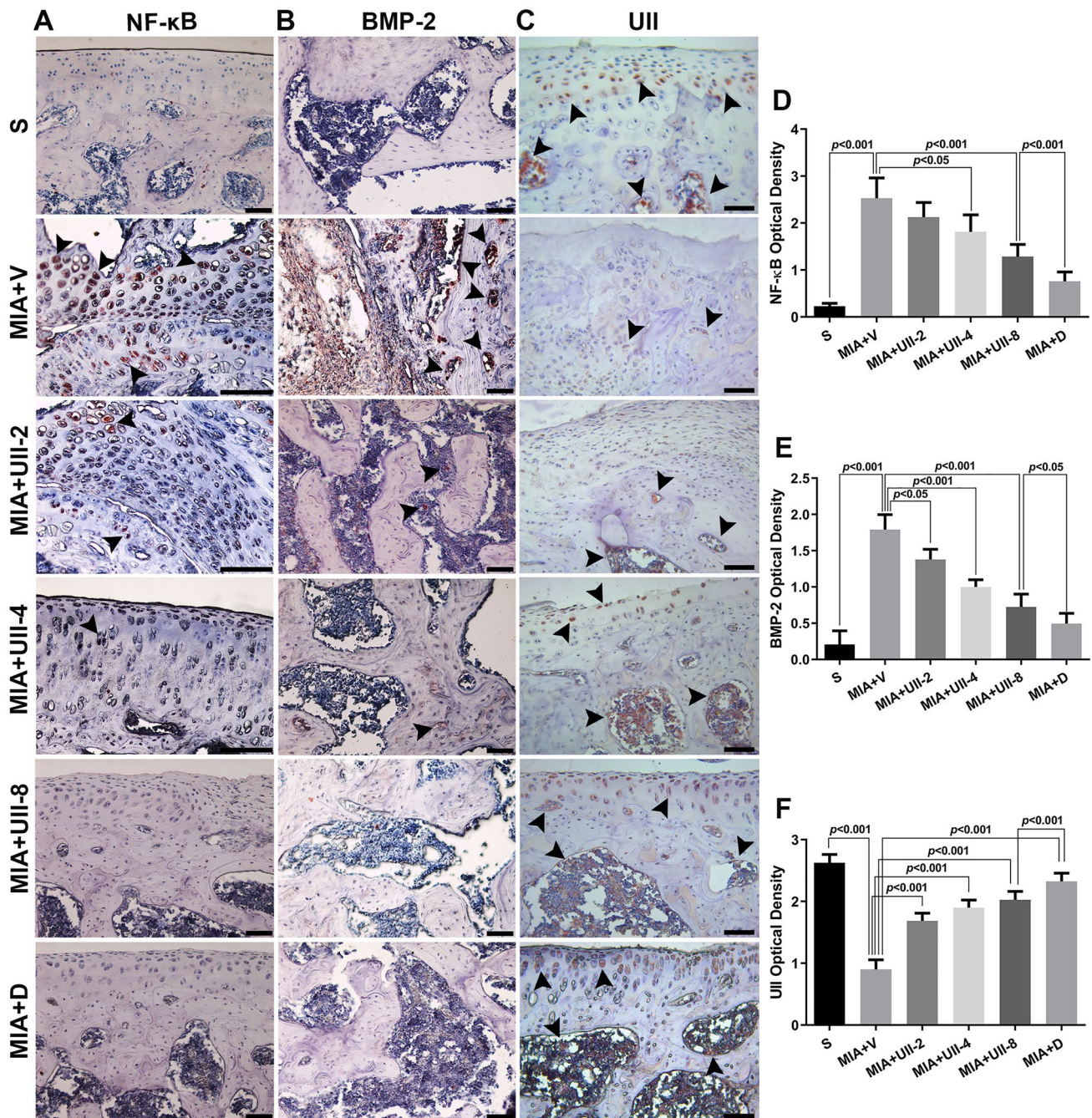


**Fig. 3** Effects of UII and diclofenac on articular cartilage histomorphology. Glycosaminoglycan and proteoglycan concentration in the OA rat model depicted with **A** H&E, **B** Toluidine blue, and **C** Safranin O & Fast Green FCF in S, MIA+V, MIA+UII-2,4,8, and MIA+D groups. Based on Mankin score grading system, plots are indicating osteoarthritic cartilage degeneration rates as **D** total Mankin score, and at **E** structure, **F** staining, and **G** cells levels sepa-

rately in S, MIA+V, MIA+UII-2,4,8, and MIA+D groups. Arrowheads point the regions with decreased cellularity in Safranin O & Fast Green FCF staining. Data were shown as mean  $\pm$  SD,  $n=8$  per group. Connected lines over bars depict significant  $p$  values between groups. Scale bar=200  $\mu$ m. *H&E* Hematoxylin and eosine, *S* Sham, *MIA* Monosodium iodoacetate, *V* Vehicle, *UII-2,4,8* Urotensin-II-2,4,8  $\mu$ g, *D* Diclofenac

biomarker in bone/cartilage pathophysiology, in the sera of osteoarthritic rats with ELISA (Shinmei et al. 1989; Kaneko et al. 2000; Braza-Boils et al. 2012; Hao et al. 2019). We found no significant alterations in the serum levels of these markers in MIA+V group compared to the

sham group (Table 2,  $p > 0.05$ ). Besides, UII or diclofenac treatments did not cause any significant change in the levels of analyzed serum markers compared to the MIA+V or sham groups (Table 2,  $p > 0.05$ ).



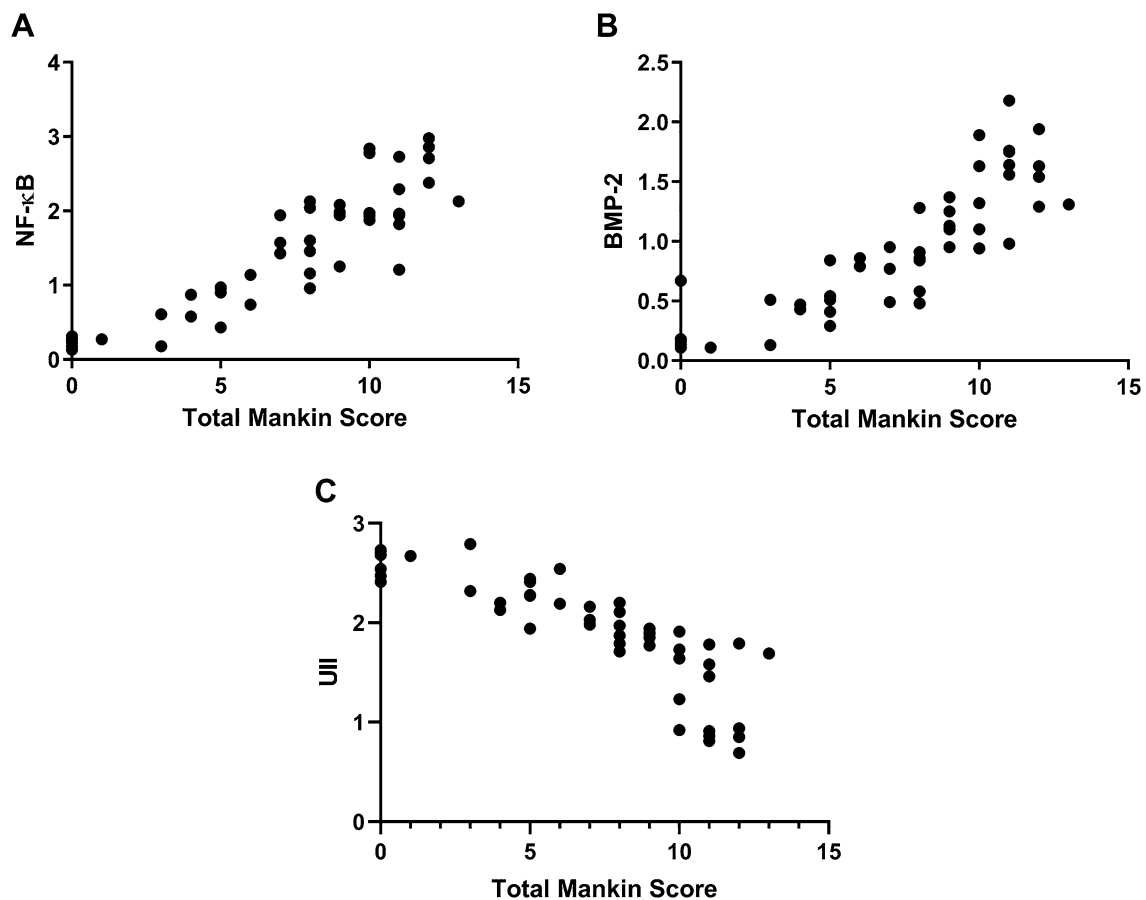
**Fig. 4** Effects of UII and diclofenac on the immunohistochemical expression of the relevant biomarkers in MIA-induced OA model in rats. Representative images of **A** NF-κB, **B** BMP-2, **C** UII stainings and **D**, **E**, **F** their corresponding expression levels (optical density) in S, MIA + V, MIA + UII-2,4,8, and MIA + D groups. Positive signals were pointed with arrowheads. Scale bar = 100 μm. Data were shown

as mean ± SD, n = 8 per group. Connected lines over bars depict significant p values between groups. *NF-κB* Nuclear factor kappa B subunit 1, *BMP-2* Bone morphogenetic protein 2, *UII* Urotensin 2, *S* Sham, *MIA* Monosodium iodoacetate, *V* Vehicle, *UII-2,4,8* Urotensin-II-2,4,8 μg, *D* Diclofenac

### Discussion

OA is a pathologic condition emerging as a result of trauma and/or aging along with several other contributory factors such as obesity, mechanical stress, and biochemical and

genetic effects (Martel-Pelletier 2004; Felson 2006). The current available therapeutic options, including surgical interventions, are mainly based on the alleviation of the pain but do not provide a full recovery of the impaired joint (Cutolo et al. 2015). The conventional medications utilized



**Fig. 5** Spearman's correlation analyses of total Mankin's scores with **A** NF- $\kappa$ B ( $r=0.8847$ ,  $p<0.05$ ), **B** BMP-2 ( $r=0.9013$ ,  $P<0.05$ ) and **C** UII ( $r=-0.9153$ ,  $p<0.0001$ ) levels in osteoarthritic joint tissues.

NF- $\kappa$ B Nuclear factor kappa B subunit 1, BMP-2 Bone morphogenetic protein 2, UII Urotensin 2

**Table 2** The effect of UII administration on serum COMP, IL-6, and IL-1 $\beta$  levels in experimental groups

Serum Parameters	S	MIA + V	MIA + UII-2	MIA + UII-4	MIA + UII-8	MIA + D	<i>p</i> value
COMP (ng/ml)	102.3 $\pm$ 6.6	93.6 $\pm$ 25.2	93.7 $\pm$ 24.8	95.0 $\pm$ 7.7	71.3 $\pm$ 21.6	95.4 $\pm$ 10.3	*/**/*** > 0.05
IL-6 (pg/ml)	64.3 $\pm$ 29.7	136.4 $\pm$ 160.3	46.6 $\pm$ 11.5	144.2 $\pm$ 169.8	187.0 $\pm$ 218.0	78.7 $\pm$ 58.1	*/**/*** > 0.05
IL-1 $\beta$ (ng/ml)	71.3 $\pm$ 26.1	75.4 $\pm$ 31.3	42.4 $\pm$ 9.1	56.4 $\pm$ 16.1	65.9 $\pm$ 29.1	51.7 $\pm$ 5.7	*/**/*** > 0.05

Data were represented as the mean  $\pm$  SD

$p > 0.05$  designates no statistical significance

\*Comparison among all groups

\*\*Comparison between MIA + V and sham groups

\*\*\*Comparison between MIA + V and MIA + UII groups

COMP Cartilage oligomeric matrix protein, IL-6 Interleukin 6, IL-1 $\beta$  Interleukin 1 beta. S sham, MIA Monosodium iodoacetate, V Vehicle, UII-2,4,8 Urotensin-II-2,4,8  $\mu$ g, D Diclofenac

in the treatment of OA can bring about serious adverse effects such as gastrointestinal tract damage and cardiovascular disorders caused by non-steroidal anti-inflammatory drugs; and cartilage damage caused by intra-articular corticosteroid injection therapies (Coxib et al. 2013; McAlindon

et al. 2017). Therefore, it is required to develop novel therapeutic drugs or strategies for long-lasting treatment of OA with minimal side effects. Thus, in the present study we sought to demonstrate the protective effects of intra-articular injection of UII on the osteoarthritic knee joint of rats. We



report for the first time that UII exerted distinct therapeutic impacts on the degenerated knee joint of rats with chemically-induced OA. We demonstrated that intra-articular UII administration significantly prevented cartilage destruction via upregulating intrinsic UII expression and downregulating NF- $\kappa$ B and BMP-2 expressions in the articular joint of osteoarthritic rats.

The studies until now have focused on the unknown onset reasons of OA and on the effective therapeutic options. In the clinical stage of the OA, the patients commonly bear inflammatory reactions ensuing synovitis which is claimed to be the primary cause of the matrix degeneration and OA (Martel-Pelletier 2004). What we know today about the pathophysiology of OA is the implication of the pro-inflammatory cytokines in articular cartilage destruction (Kim et al. 2018). The first clear histopathological symptom of the OA is diminishing of the articular collagen and proteoglycan structures due to activated pro-inflammatory cytokines which eventually leads to degradation of synovial matrix via several matrix metalloproteases and aggrecanases (Tortorella et al. 1999; Braza-Boils et al. 2012). In a MIA-induced OA rat model, the cartilage degeneration was observed related to the time-dependent loss of proteoglycan anabolism, increased gelatinase activity, and IL-1 $\beta$ -mediated inflammatory response in articular regions (Dumond et al. 2004). The previous clinical findings support the claim that elevated cytokine levels such as IL-1 $\beta$ , TNF- $\alpha$ , and IL-6 can cause inflammation-induced cartilage degeneration in OA patients (Shinmei et al. 1989; Dumond et al. 2004). Supporting this, the SF levels of inflammatory cytokines such as IL-1 $\beta$ , IL-6, IL-8, etc. significantly increased in OA patients (Kim et al. 2018). In our study, we did not find any significant alteration in the serum levels of COMP, IL-6, and IL-1 $\beta$  in OA-induced rats. A study conducted with the spontaneous OA model of mice indicated that OA onset and progression did not affect serum levels of IL-1 $\beta$ , TNF- $\alpha$ , and COMP (Braza-Boils et al. 2012). This finding is in line with the steady IL-1 $\beta$ , IL-6, and COMP levels after OA induction in our study. It may be partially due to the fact that the alterations in articular OA-related biomarkers are not always correlated with their corresponding systemic serum levels (Kaneko et al. 2000).

As aforementioned above, UII is a short cyclic vasoactive peptide with a known expression pattern in several tissues of the human body except knee joints (Pearson et al. 1980). After immunohistochemical analyses, our group was to be the first to demonstrate the articular expression pattern of UII in knee joints of the rats. Firstly, MIA-induction significantly lowered the UII expression in osteoarthritic tissue of the rats compared to the sham group. After UII treatment, we found a significant increase in articular UII expression in a dose-dependent manner. It was claimed that there is a close connection between UII expression and the inflammatory

reactions in onset or progression of several diseases (Gogebakan et al. 2014; Sun and Liu 2019). In our OA model, however, we found that intra-articular UII injection incited the intrinsic expression of UII. Furthermore, there was a strong negative correlation between articular UII expression and tissue degeneration level. It can be inferred from our findings that the restoration of intrinsic UII expression after extrinsic UII administration in the articular region prevented the further MIA-induced destruction of cartilage tissue in the rats.

Several signaling pathways have been shown to be implicated in the pathogenesis of OA including NF- $\kappa$ B (Kim et al. 2018). NF- $\kappa$ B acts as a central molecule in promoting inflammation-induced cartilage degeneration in OA by igniting the inflammatory cytokines, which can re-stimulate NF- $\kappa$ B in a positive feedback manner (Rigoglou and Papatavassiliou 2013). The NF- $\kappa$ B pathway was also implicated in chondrocyte apoptosis in an IL-1 $\beta$ -induced cell line model (Pan et al. 2018). Supporting the idea, it was demonstrated that the knock-down of NF- $\kappa$ B ameliorated the cartilage breakdown in a surgical rat model of OA (Chen et al. 2008). Along with the previous findings, we showed the downregulation of NF- $\kappa$ B expression in knee joint after UII treatment that contributed to the slowdown of the osteoarthritic knee joint degeneration. On the other hand, the blockade of UII/UTR via its specific antagonist was previously shown to attenuate the NF- $\kappa$ B-activated inflammatory reactions in *in vivo* and *in vitro* inflammatory-disease models (Liang et al. 2014; Liu et al. 2015a, b). Another *in vivo* sepsis-induced lung injury animal model demonstrated the upregulation of NF- $\kappa$ B levels after intraperitoneal UII administration (Ugan et al. 2018). Our results seem to be contradicting with these findings; however, we administered exogenous UII peptide locally in our model rather than systemic injection or intrinsic manipulation of UII expression and/or UII/UTR pathway via specific inhibitors. Our strategy could likely interfere and distort the intrinsic action mechanism of UII/URT pathway owing to the paracrine and autocrine natures of UII activity over intra- and extra-cellular regions (Castel et al. 2017). As aforementioned, several pathways are implicated in OA progression like apoptosis. In an *in vitro* oxidative stress cardiomyocyte cell line model, UII treatment reduced apoptosis via ERK upregulation (Gong et al. 2015) which presumably down modulated the NF- $\kappa$ B-induced apoptosis (Shi et al. 2015). In our model, the downregulation of NF- $\kappa$ B post-UII treatment can be implied as the protection of articular cells from apoptosis induced by OA.

NF- $\kappa$ B pathway is mostly triggered together with other pathways including BMP to inhibit chondrogenesis and activate other matrix-degrading enzymes (Lepetosos et al. 2019). Several lines of evidence revealed that BMP pathway-related proteins can play a central role in the formation of hypertrophic chondrocytes that causes matrix degeneration during

OA pathogenesis via upregulation of matrix metalloproteinases (van der Kraan et al. 2010; Brescia et al. 2014). Moreover, it was demonstrated that NF- $\kappa$ B could also stimulate the upregulation of BMP-2 in hypertrophic chondrocytes in a transgenic mouse model (Ijiri et al. 2005). The upregulation of BMP-2 expression at gene levels was also shown in an exercise-induced OA rat model (Liu et al. 2016). On the one hand, the anabolic and chondroprotective effects of BMP family proteins are also known (Liu et al. 2015a, b); thus the significant upregulation of BMP-2 within the osteoarthritic cartilage region in our model can also be an endogenous chondroprotective response to cartilage degeneration. On the other hand, articular BMP-2 administration has been previously demonstrated to cause osteophytes in the ectopic regions of murine knee cartilage (Scharstuhl et al. 2003). We also demonstrated the significant reduction in BMP-2 expression ensuing tissue recovery concurrently in UII-treated tissues, which clearly exhibits the involvement of BMP-2 and UII in articular tissue physiology. In line with our data, the central role of the BMP-2 pathway has been shown in both in vitro and in vivo models of OA by suggesting that IL-1 $\beta$  increased BMP-2 expression and blocking of BMP-2 via its inhibitors improved the degenerated cartilage tissue in knee joints of rats with OA (Chien et al. 2020). It was also previously suggested that transforming growth factor-beta 1/BMP signaling pathways are of critical importance for the maintenance of cartilage homeostasis, and any disturbance of this balance in favor of BMP can result in cartilage chondrocyte loss (Shen et al. 2017). Moreover, we have found significant positive correlations between histopathological lesion rate and tissue-specific expressions of NF- $\kappa$ B and BMP-2 pointing out the fact that both proteins are valid markers in assessing cartilage degeneration.

As to the best of our knowledge, we are the first group investigating the effect of articular UII treatment on the pathogenesis of our MIA-induced OA model in rats. We demonstrated that UII treatment significantly restrained the MIA-induced articular tissue degeneration and partially protected the cartilage and subchondral bone as depicted in gross morphology and histological and immunohistochemical analyses (Figs. 2, 3, 4). In previous studies, UII has been linked to the pathogenesis of several diseases with respect to inflammatory responses (McDonald et al. 2007). A clinical study conducted with OA and non-OA patients suggested that the elevated levels of UII in the SF of the OA patients can be related to the synovial fibrosis during OA progression (Gogebakan et al. 2014). It has been also reported decades ago that UII acts as an osmoregulator to establish the hemostasis of the body fluids not only in fishes but also in vertebrates, including humans (Balment et al. 2005). The SF is produced by ultrafiltration from blood plasma by synovial cells. Considering the fact that the deterioration of the SF quality and quantity during OA

pathogenesis (More et al. 2020; Seidman and Limaïem 2020), we can assume that the elevation of endogenous synovial UII expression in the patients with knee OA can be a protective response against cartilage destruction due to the reduced SF in both quality and quantity. Supporting this idea, a previous study demonstrated the expression of UII receptor in the chondrocytes within hyaline cartilage of amphibians and rats and presumed a modulatory role of UII in cartilage matrix remodeling (Konno et al. 2013).

Last but not least, in our study, the exogenous intra-articular UII administration could prevent the inflammation-induced destruction of articular matrix composition presumably by interfering the endogenous UII/UTR pathway regulated by autocrine and paracrine mechanisms as ascertained previously (Castel et al. 2017). However, the protective effect of UII in the present model can be context-specific, meaning that extrinsic UII can act differently in other in vivo systems e.g., surgical or age-induced OA models.

## Conclusions

Based on our findings, the external UII treatment could slow down cartilage degeneration and help to prevent the cartilage tissue damage through downregulation of NF- $\kappa$ B and BMP-2 proteins and compensation of the intrinsic UII expression via inducing its upregulation. Since UII is a strong vasoactive peptide, it can be speculated that the restored intrinsic UII expression in the joint tissue may reconstitute the rheological characteristics of the synovial fluid load. To the best of our knowledge, there are no studies in the literature analyzing the effect of exogenous UII administration on the knee joint of MIA-induced osteoarthritic rats. To elucidate the underlying action mechanism of UII on knee joint pathophysiology during OA, further experiments are warranted.

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## Declarations

**Conflict of interest** There are no conflicting interests.

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## Authors and Affiliations

Menderes Yusuf Terzi<sup>1,2</sup>  · Hamza Malik Okuyan<sup>3</sup> · İhsan Karaboğa<sup>4</sup> · Cemil Emre Gökdemir<sup>5</sup> · Duygu Tap<sup>2</sup> · Aydiner Kalaci<sup>5</sup>

✉ Menderes Yusuf Terzi  
menderesyusufterzi@gmail.com; myterzi@mku.edu.tr

<sup>1</sup> Department of Medical Biology, Faculty of Medicine, Hatay Mustafa Kemal University, Hatay, Turkey

<sup>2</sup> Department of Molecular Biochemistry and Genetics, Graduate School of Health Sciences, Hatay Mustafa Kemal University, Hatay, Turkey

<sup>3</sup> Department of Biomedical Engineering, Department of Physiotherapy and Rehabilitation-Faculty of Health

Sciences, Sakarya University of Applied Sciences, Sakarya, Turkey

<sup>4</sup> Department of Emergency and Disaster Management, School of Health, Tekirdağ Namık Kemal University, Tekirdağ, Turkey

<sup>5</sup> Department of Orthopedics and Traumatology, Faculty of Medicine, Hatay Mustafa Kemal University, Hatay, Turkey