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Erythromycin acts through the ghrelin receptor to attenuate inflammatory responses in chondrocytes and maintain joint integrity

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Abstract

Osteoarthritis (OA) is a prevalent disease characterized by chronic joint degeneration and low-grade localized inflammation. There is no available treatment to delay OA progression. We report that in human primary articular chondrocytes, erythromycin, a well-known macrolide antibiotic, had the ability to inhibit pro-inflammatory cytokine Interleukin 1 β (IL-1 β)-induced catabolic gene expression and nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B) activation. Furthermore, erythromycin inhibited monosodium iodoacetate (MIA) -induced joint inflammation and cartilage matrix destruction in mice, an arthritis model that reflects the inflammatory and cartilage matrix loss aspects of OA. EM900, an erythromycin-derivative lacking antibiotic function, had the same activity as erythromycin *in vitro* and *in vivo*, indicating distinct anti-inflammatory and antibiotic properties. Using an antibody against erythromycin, we found erythromycin was present on chondrocytes in a dose-dependent manner. The association of erythromycin with chondrocytes was diminished in ghrelin receptor null chondrocytes, and administration of the ghrelin ligand prevented the association of erythromycin with chondrocytes.

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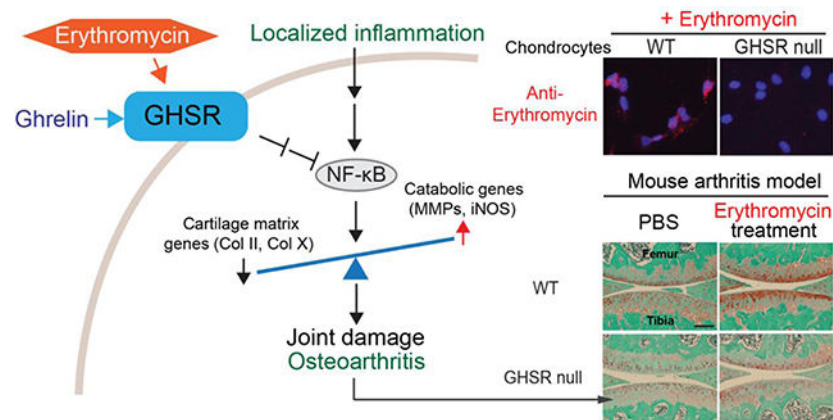
Conflict of interest

None of the authors have any conflict of interest related to this work.

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Importantly, the anti-inflammatory activity of erythromycin was diminished in ghrelin receptor null chondrocytes. Moreover, erythromycin could not exert its chondroprotective effect in ghrelin receptor null mice, and the loss of ghrelin receptor further augmented joint damage upon MIA-injection. Therefore, our study identified a novel pharmacological mechanism for how erythromycin exerts its chondroprotective effect. This mechanism entails ghrelin receptor signaling, which is necessary for alleviating inflammation and joint destruction.

Graphical Abstract



Keywords

Erythromycin; ghrelin receptor; osteoarthritis; chondrocyte; cartilage; synovitis

1. Introduction

Osteoarthritis (OA) is a prevalent disease characterized by chronic joint degeneration, including articular cartilage loss, synovitis and osteophyte formation [1–5]. The cause of OA is metabolic imbalance and mechanical stress, which is coupled with inflammation, leading to a catabolic shift in cartilage homeostasis. Increased macrophage activity has been reported as a biomarker for human OA [6], and antagonizing pro-inflammatory cytokine Interleukin 1 β (IL-1 β) directly in the joint protects cartilage matrix in post-traumatic experimental OA, indicating a pivotal role of inflammation in OA pathogenesis [7]. IL-1 β activity is mediated by the nuclear factor kappa-light-chain-enhancer of activated B cells (NF- κ B) signaling pathway, which upon activation induces the expression of cartilage matrix-degrading enzymes such as matrix metalloproteinases (MMPs) and subsequent matrix degradation [8].

Recently, certain antibiotics used to treat bacterial infections have been shown to alter the response of chondrocytes toward inflammatory stimuli. Tetracycline and doxycycline chelate zinc at the active sites of MMPs and inhibit experimental OA and rheumatoid arthritis [9–14]; however, reduction in human joint pain and swelling was limited [15, 16]. The macrolide class of antibiotics, including erythromycin (EM) and azithromycin, has also been found to be anti-inflammatory. In patients with diffuse panbronchiolitis, they inhibit

macrophage activation and inflammation in the lung [17–19]. Similarly, EM inhibited aseptic loosening after bone implants [20, 21]. Our group recently showed that EM inhibits inflammatory stimuli IL-1 β and Lipopolysaccharides (LPS)-induced catabolic events in bovine articular chondrocytes [22]. However, the mechanism underlying this activity remains poorly understood.

In the current study, we discovered that the activity of EM is mediated by the ghrelin receptor. Ghrelin is a 28 amino acid peptide that is best known as the “hunger hormone,” as it increases appetite and controls blood sugar levels [23–26]. The ghrelin receptor is also known as the Growth Hormone Secretagogue Receptor (GHSR1a, or abbreviated as GHSR) [27]. While a truncated variant GHSR1b also exists, it does not bind to ghrelin; therefore, ghrelin signals through its binding to GHSR1a [28]. In chondrocytes, ghrelin has been found to control fatty acid uptake and inhibit inflammation when overexpressed [29–31]. However, these are gain-of-function experiments, and it is still not known whether ghrelin signaling is truly necessary for joint maintenance under disease conditions.

We identified a novel functional link between EM and ghrelin signaling in chondrocytes, since the loss of the ghrelin receptor abolished EM-mediated chondro-protection and resulted in more cartilage destruction *in vivo*. Administration of ghrelin to chondrocytes had a similar inhibitory activity as EM toward pro-inflammatory cytokine IL-1 β -induced catabolic gene expression. This activity was diminished in ghrelin receptor null cells, suggesting that EM and ghrelin both act through the ghrelin receptor.

2. Materials and Methods

2.1. Experimental animals

All animal care and experimental procedures were approved by the Institutional Animal Care and Use Committees at Tufts University. C57/B6 mice were used. The ghrelin receptor null mouse (i.e. GHSR null, in C57/B6 background) was a generous gift from Dr. Jeffrey Zigman, University of Texas Southwestern Medical Center [26]. Mice were caged in groups under the standard conditions with a 12-hour light/dark cycle, fed with a standard chow diet and identified by ear tags. Tail biopsies were used for PCR-based genotyping based on the established method with primers (5' to 3'): Forward: CCACTGCACGTCTCTCCCTATTT and reverse for WT: CGGTCTCCACCCTTCATTACTTTA or Rev for null: AAGAGCTACAGGAAGGCAGGTCA [26]. Mice were euthanized by carbon dioxide overdose followed by cervical translocation or by decapitation under inhalation anesthesia.

2.2. Isolation of normal and OA human cartilage specimens for RT-PCR analysis

Normal and OA human cartilage specimens were obtained as previously described [32], with additional samples over time. Samples of severe OA that showed no presence of articular cartilage were excluded. Articular cartilage slices of normal and OA specimens were excised from the tibial plateau of cadaveric joints (National Disease Research Interchange and Articular Engineering, age/sex: 84/M, 75/M, 65/F, 49/F). Human OA tibial plateau was obtained from patients undergoing total knee replacement surgery at Tufts Medical Center (age/sex: 53/F, 63/F, 65/F, 73/F, 72/M). All cartilage samples were de-identified before we

received them and would have otherwise been discarded. Sample collection protocols were reviewed by the Institutional Review Board (IRB) at Tufts University and classified as exempt. Cartilage explants from cadavers were refrigerated for no longer than 36hrs postmortem before being used for chondrocyte isolation or gene expression analysis. Cartilage from knee replacement surgeries were obtained within 1hr after the excision. Due to the avascular nature of articular cartilage, chondrocytes in excised cartilage specimens remain alive for long periods of time, and have been used postmortem for culturing and gene expression analysis [33–37]. Each specimen was divided into two groups. The first group was subjected to histological analysis to confirm the integrity of the articular cartilage as previously reported [32]. The second group was subjected to gene expression analysis by RT-PCR.

2.3. In vitro culture of primary normal human articular chondrocytes

Primary normal human articular chondrocytes (nHACs) were purchased from Lonza (Walkersville, MD), or isolated from cadaveric joints retrieved within 36hrs post-mortem (National Disease Research Interchange). Experiments were performed using chondrocytes from three donors: age/sex: 49/M, 38/M, and 84/M). Before use, nHACs were re-differentiated per Lonza's instructions. Briefly, nHACs were encapsulated in 1.2% alginate beads (Sigma, St. Louis, MO) at a density of 8×10^5 cells/ml and cultured in chondrocyte differentiation media (CDM) containing TGF- β 1 (Lonza) supplemented with ascorbic acid for 2 to 4 weeks. Alginate beads were dissolved in 55mM citric acid (Sigma) supplemented with 10mM HEPES (Thermo Fisher Scientific, Waltham, MA) to collect re-differentiated nHACs. Cell viability was confirmed by 3-[4,5-dimethylthiazole-2-yl]-2,5-diphenyltetrazolium bromide (MTT) assay (Sigma). Usually more than 80% cell viability was obtained. Re-differentiation of chondrocytes was confirmed by elevated Collagen II (Col-II) mRNA expression and alcian blue (Thermo Fisher Scientific) staining before proceeding. Once re-differentiation was confirmed, cells were treated with EM (Sigma), EM900 (a 12-membered macrolide and EM derivative manufactured in Drs. Sunazuka and Ömura laboratories) and ghrelin (Phoenix Pharmaceuticals, Burlingame, CA) overnight, and then subjected to IL-1 β (1ng/ml) (Peprotech, Rocky Hill, NJ) treatment. For RT-qPCR analysis of EM, EM900 and ghrelin-treatment, cells were lysed after 3 days of IL-1 β treatment. For RT-qPCR analysis of MIA-treated chondrocytes *in vitro*, cells were analyzed after 2 days of MIA (Thermo Fisher Scientific) treatment. All experiments were performed at least three times.

2.4. In vitro culture of murine articular chondrocytes

Murine femoral and tibial chondrocytes were isolated based on a previously published protocol [38]. Briefly, cartilage pieces from the femoral head, femoral condyle and tibial plateau were isolated and digested in collagenase D (Thermo Fisher Scientific) to remove residual soft tissues and to isolate chondrocytes. WT and GHSR null chondrocytes at passage 1 were seeded in a well of a 24-well plate in DMEM/F-12 (Thermo Fisher Scientific) supplemented with 10% FBS (Thermo Fisher Scientific). For analyzing the presence of EM in chondrocytes, cells were seeded in a well of an 8-wells chamber (Thermo Fisher Scientific) before being treated with EM (Sigma) or co-treated with ghrelin (Phoenix Pharmaceuticals) in serum-free media for 1 hour. For RT-qPCR analysis, cells were seeded

in wells of a 24-wells plate. After 24 hours, EM or ghrelin were added to the cells, followed by IL-1 β (1ng/ml) (Peprotech) treatment on the next day for an additional 3 days. All experiments were performed at least three times.

2.5. Explant culture of human articular cartilage

For *ex vivo* culturing of OA cartilage slices, OA cartilage pieces were prepared from OA donors as described above (72/M, 69/M, 73/F, 67/M, 54/F, 53/F, 60/F). Cartilage slices were cut into 2 \times 3 \times 2mm and cultured in the presence or absence of ghrelin (100nM), EM or EM900 (10 μ M) for 3 weeks with media changes every 3 days. Explant cultures were performed in duplicate per group and repeated for at least three donors.

2.6. Monosodium iodoacetate (MIA)-induced articular cartilage degeneration model

Adult male mice (18–22 weeks) were anesthetized by isoflurane/O₂ inhalation. EM or EM900 at 40 μ g/g body weight/day was delivered by intraperitoneal injection using a 28G $\frac{1}{2}$ " needle. PBS was delivered as a control. 5 μ L of 10 μ g/ μ L MIA dissolved in PBS was injected intra-articularly using a 30G $\frac{1}{2}$ " needle. PBS was injected to contralateral knees as a control. EM or EM900 was administered every day until mice were euthanized at 7 days post injection of MIA. For each experiment, at least n=4 mice were used.

2.7. Cytotoxicity assay of erythromycin (EM) and EM900

Chondrocytes were seeded on tissue culture dishes and treated with EM and EM900 (0–100 μ M) for 4 days before being subjected to Live/Dead Cell Viability/Cytotoxicity assay (Thermo Fisher Scientific) according to the manufacturer's instruction. The ratios of dead cells/total number of cells were determined.

2.8. RT-qPCR

Total RNA was isolated using the RNeasy mini kit (Qiagen, Germantown, MD) and reverse transcribed using the M-MLV reverse transcriptase (Thermo Fisher Scientific). Quantitative PCR was carried out using iTaq universal SYBR Green supermix (Bio-Rad, Hercules, CA) on the iQ5 Real Time Detection System (Bio-Rad). TATA-binding protein (TBP) served as a reference gene. The sequences for PCR primers are: human primers: TBP, forward 5'-GGTGCTAAAGTCAGAGCAGAA-3', reverse 5'-CAAGGGTACATGAGAGCCATTA-3'; Ghrelin, for, 5'-AAGTGATCGCCACAAGCCTTACT-3', reverse, 5'-TGTACAACAGTCGTGGGAGTTGCT-3'; Aggrecan (ACAN), forward, 5'-AAGGTCTCTCTCTCAGCCACCT-3', reverse, 5'-TGGCTGAGGGAATGGACACTGAAA-3'; Collagen II (COL-II), forward, 5'-TTCATCCCACCCTCTCACAGTT-3', reverse, 5'-CCTCTGCCTTGACCCGAA-3'; Collagen IX (COL-IX), forward, 5'-TCGATGTGCTGTCTCTGGAGTGAT-3', reverse, 5'-CTTTGCTTTTCACATAGAAGCGCTCA-3'; Collagen X (COL-X), forward, 5'-ACCCAGCATAACTTGAAACAGGT-3', reverse, 5'-TCACTTGAATGGGAGGCACAAGGT-3'; IL-1 β , forward, 5'-AACAGGCTGCTCTGGGATTCTCTT-3', reverse, 5'-ATTTCACTGGCGAGCTCAGGTACT-3'; TNF α (tumor necrosis factor alpha), forward, 5'-AGGACGAACATCCAACCTTCCCAA-3', reverse, 5'-

TTTGAGCCAGAAGAGGTTGAGGGT-3'; iNOS (inducible nitric oxide synthase), forward, 5'-GCGTTACTCCACCAACAATGGCAA-3', reverse, 5'-ATAGCGGATGAGCTGAGCATTCCA-3'; MMP1, forward, 5'-AGTGACTGGGAAACCAGATGCTGA-3', reverse, 5'-TCAGTGAGGACAAACTGAGCCACA-3'; MMP3, forward, 5'-AGGCAAGACAGCAAGGCATAGAGA-3', reverse, 5'-ACAAGGTTTCATGCTGGTGTCTCA-3'; MMP9, forward, 5'-GACCTGGGCAGATTCCAAA-3', reverse, 5'-GGCAAGTCTTCCGAGTAGTTT-3'; MMP13, forward, 5'-ACAAGTAGTTCCAAAGGCTACAA-3', reverse, 5'-ATAGGAAACATGAGTGCTCCAG-3'. Mouse primers: TBP, forward, 5'-CTACCGTGAATCTTGGCTGTAA-3', reverse, 5'-GTTGTCCGTGGCTCTCTTATT-3'; MMP13, forward, 5'-TCTGGGCTCTGAATGCTTATG-3', reverse, 5'-GCTCAGTCTCTTACCTCTTT-3'.

2.9. NF- κ B transactivation assay

70ng of a six-copy NF- κ B element-driven luciferase reporter construct and 10ng of Renilla luciferase internal control were transiently co-transfected into nHACs using X-tremeGENE HP (Roche, Indianapolis, IN). EM (10 μ M), EM900 (10 μ M) or ghrelin (10nM) was added at the time of transfection. At 24 hours after the transfection, chondrocytes were treated with 1ng/ml IL-1 β for an additional 16 hours. Luciferase assay was conducted using the Dual Luciferase Assay kit (Promega, Madison, WI). Luminescence was detected using a microplate luminometer (Perkin Elmer, 1450 Microbeta TriLux). NF- κ B luciferase activity was normalized to Renilla luciferase activity. Fold induction of normalized luciferase activity was determined based on the control sample as a baseline. At least three independent experiments in triplicate were performed.

2.10. Serum response element (SRE) reporter assay

100ng of a SRE luciferase reporter construct (gift from Dr. Alan Kopin, Tufts Medical Center) and 30ng of Renilla luciferase internal control were transiently co-transfected into nHACs using X-tremeGENE HP (Roche). At 24 hours after the transfection, cells were treated with EM (10 μ M), EM900 (10 μ M) or ghrelin (10nM) in serum-free media for additional 18 hours. Luciferase assay was conducted using the Dual Luciferase Assay kit (Promega). Luminescence was detected using a microplate luminometer (Perkin Elmer, 1450 Microbeta TriLux). SRE luciferase activity was normalized to Renilla luciferase activity. Fold induction of normalized luciferase activity was determined based on the control sample as a baseline. At least three independent experiments in triplicate were performed.

2.11. Immunocytochemistry (ICC) and histological analysis

For ICC analysis, chondrocytes were fixed with ice-chilled ethanol (VWR, Radnor, PA) for 15 minutes. Cells were blocked with goat serum (Thermo Fisher Scientific) for 1 hour and incubated with primary sheep anti-EM antibody (Abcam, ab123983, Cambridge, MA) followed by incubation with Alexa594 or fluorescein-conjugated secondary goat anti-sheep antibody (Thermo Fisher Scientific). Nuclei were visualized by DAPI (4',6-diamidino-2-phenylindole, Thermo Fisher Scientific) staining. Five different views under the microscope

were randomly selected from each well for quantification. Experiments were performed in triplicates.

For histological analysis of joint tissues, samples were fixed in 1% PFA (VWR, Radnor, PA), then decalcified in 0.33M EDTA (VWR) 10 days for mouse knee joints and overnight for human cartilage specimens. Decalcified samples were dehydrated in 25, 50, 75, 100 and 100% ethanol/PBS and xylene twice for each 1hr. Samples were embedded in paraffin (Paraplast plus, McCormick Scientific, St. Louis, MO). Mouse knee joints and human cartilage specimens were sectioned at 5 μ m thickness.

Sections were stained with 0.1% Safranin O (Sigma) and counterstained with Hematoxylin (Sigma) and Fast green (Sigma). Cartilage integrity was evaluated by the established OARSI scoring system that is based on loss of Safranin O staining, as described previously [32, 39]. Severity of synovitis was scored by the established method [40, 41]. This scoring system has the following three criteria: enlargement of the synovial cell layer (0–3 points), cellularity of resident cells (0–3 points) and inflammatory infiltration (0–3 points). When the points are totaled, a full score of 9 points would indicate severe synovitis with lower scores indicating less severe synovitis. Three to four sections per mouse knee joint or human cartilage slices within 100 μ m intervals were evaluated and scored. All scorings were performed blinded.

For ghrelin receptor (GHSR) immunohistochemistry, heat-induced antigen retrieval in 10mM citric acid buffer at pH 6.0 was performed for 10min by steam. Primary GHS-R1a antibody (Santa Cruz, F-16, Dallas, Texas) and secondary biotin-conjugated horse anti-goat antibody (Vector labs, Burlingame, CA) were used. For chromogenic staining, the Vectastain ABC Elite kit (VWR, Radnor, PA) was used according to the manufacture's instruction. Nuclei were counterstained with methyl green (Sigma).

2.12. Glucose consumption assay

Glucose consumption assays were performed as done previously [42]. Cells were seeded overnight and then treated with MIA for 2 days. OA and nHACs at passage 1 were seeded and grown for 2 days. Conditioned media were harvested at the end of culture and glucose concentration was determined using Glucose (HK) assay kit (Sigma) according to the manufacturer's instruction. Briefly, 3 μ L of media samples and glucose standards were incubated with 250 μ L assay reagent for 15 minutes and the absorbance at 340nm was measured using a plate reader (Bio-Rad Benchmark Plus). Total glucose consumption amount was calculated by subtracting the total glucose amount of each sample from the initial glucose amount. Total cell numbers were used to for normalization. For each treatment, technical duplicates were performed, with triplicate biological repeats.

2.13. Intracellular ATP assay

Intracellular ATP assays were performed as done previously [42]. Cells were seeded overnight and then treated with MIA for 2 days. OA and nHACs at passage 1 were seeded and grown for 2 days. At the end of culture, cells were lysed in the lysis buffer (200mM Tris pH.7.5, 2M NaCl, 20mM EDTA and 0.2% Triton X-100) for 5 minutes on ice, and supernatants were collected for ATP assay using ATP Determination kit (Thermo Fisher Scientific) according to the manufacturer's instruction. Each sample and ATP standards were

incubated with reaction solutions for 2 minutes and luminescence was detected using a microplate luminometer (Perkin Elmer, 1450 Microbeta TriLux). Total cell numbers were used for normalization. For each treatment, technical duplicates were performed, with triplicate biological repeats.

2.14. Reactive oxygen species (ROS) assay

ROS assays were performed as done previously [42]. Cells were seeded overnight and then treated with MIA for 2 days. OA and nHACs at passage 1 were seeded and grown for 2 days. ROS was detected using Image-iT Live Green Reactive Oxygen Species Detection Kit (Thermo Fisher Scientific) according to the manufacturer's instruction. Cells were washed with HBSS/Ca/Mg and incubated with 25 μ M carboxy-H₂DCFDA (2',7'-Dichlorodihydrofluorescein diacetate) for 30 minutes at 37°C. ROS production was measured at 495/529nm (excitation/emission) using a plate reader (FlexStation 3, Molecular devices). ROS production was normalized by signal of Hoechst 33342 measured at 350/461nm (excitation/emission). For each treatment, technical duplicates were performed, with triplicate biological repeats.

2.15. Light microscopy

Bright field images were taken using an Olympus IX-71 microscope and Olympus DP70 digital cameras. The optical parameters and camera exposure time were kept constant between samples of the same experiment.

2.16. Statistical analysis

Parametric data were reported as mean \pm standard deviation (SD) and statistical analysis was performed using a Student's *t*-test, Dunnett's test or a one-way analysis of variance followed by a post-hoc Tukey test with *p*-values of <0.05 considered significant. Non-parametric data (OARSI scores and synovitis scores) were reported as a box plot, where the median, the maximum and minimum data values were indicated; and statistical analysis was performed using the Kruskal-Wallis test followed by Mann-Whitney *U* test with *p*-values of <0.05 considered significant. All statistical analyses were conducted on Prism (Graphpad).

3. Results

3.1. EM inhibits pro-inflammatory cytokine IL-1 β -induced catabolic gene expression and NF- κ B activity in chondrocytes

To evaluate EM activity on human articular chondrocytes under inflammatory conditions, we treated primary normal human articular chondrocytes (nHACs) with EM and IL-1 β . A "live-dead" assay was first performed on nHACs to determine the concentrations of EM to be used. We tested 1, 10 and 100 μ M of EM, which are in the range of systemic plasma concentrations achieved with typical therapeutic activity of EM in humans [43]. The level of cell death overall was minimal (1–4%), although higher EM concentrations caused slightly more cell death (Fig. 1A). As expected, IL-1 β reduced cartilage matrix collagen II (Col-II) mRNA expression, and induced hypertrophic marker collagen X (Col-X), as well as catabolic genes iNOS and MMPs expression (Fig. 1B). EM treatment alone did not alter the expression of these genes. However, when chondrocytes were pre-treated with EM, the

induction of hypertrophic and catabolic genes Col-X, iNOS and MMPs by IL-1 β was strongly reduced, and Col-II expression restored (Fig. 1B). This result suggests that EM itself inhibits the response to pro-inflammatory stimuli in human chondrocytes.

To further investigate the underlying mechanism, we evaluated activation of NF- κ B, the key mediator of IL-1 β signaling [32, 44]. nHACs were transfected with an NF- κ B luciferase reporter, which has NF- κ B binding sites and serves as a readout for NF- κ B transcriptional activity after NF- κ B nuclear translocation [32, 45] (Fig. 1C). EM alone did not have any effects on the activity of this luciferase reporter construct. In contrast, IL-1 β clearly induced NF- κ B reporter activity, but co-treatment with EM significantly reduced IL-1 β -induced luciferase activity (Fig. 1C). Thus, EM inhibits the ability of chondrocytes to activate NF- κ B signaling and induce catabolic gene expression under inflammatory conditions.

3.2. Ghrelin receptor (GHSR) mediates the activity of erythromycin (EM) to antagonize IL-1 β

To further understand EM activity on chondrocytes, we searched the literature for proteins that interact with EM. Interestingly, classic *in vitro* binding assays showed that a derivative of EM could bind to the ghrelin receptor [46–48] and ghrelin signaling inhibited inflammation [30, 31]. Thus, we hypothesized that ghrelin signaling might mediate the activity of EM. This hypothesis was tested through the use of the mouse strain lacking the ghrelin receptor GHSR, i.e. GHSR null mice. Immunohistochemistry (IHC) analysis confirmed that GHSR is expressed in articular chondrocytes and is absent in the GHSR null mice (Fig. 2A). Subsequently, primary chondrocytes from WT and GHSR null mice were used to determine whether EM requires GHSR to inhibit IL-1 β activity. EM was capable of inhibiting IL-1 β -induced MMP13 expression in WT cells, but not in GHSR null cells (Fig. 2B), suggesting that EM requires the ghrelin receptor for this activity. Next, we evaluated whether GHSR mediates the activity of EM on chondrocytes. In our immunocytochemistry analysis, EM was associated with chondrocytes in a dose-dependent manner (Fig. 2C). Furthermore, this binding was diminished in chondrocytes from the GHSR null mice (Fig. 2D), suggesting that the association of EM to chondrocytes is mediated by the GHSR. However, it is not clear whether EM and ghrelin compete for the same site on the ghrelin receptor.

3.3. Ghrelin exhibits a similar activity as EM for inhibiting IL-1 β -induced catabolic gene expression and NF- κ B activation

Since we found that the ghrelin receptor mediates the activity of EM on chondrocytes under IL1 β treatment, we further hypothesized that the natural ligand for the ghrelin receptor, ghrelin, would have a similar activity. Therefore, normal primary human articular chondrocytes (nHACs) were treated with IL-1 β in the absence or presence of ghrelin and the response to IL-1 β was assayed. While ghrelin treatment did not reduce IL-1 β -induced downregulation of Col-II mRNA expression, it inhibited IL-1 β -induced Col-X expression as well as those of catabolic genes iNOS and MMPs (Fig. 3A). In addition, ghrelin strongly inhibited IL-1 β -induced NF- κ B luciferase activity (Fig. 3B). Interestingly, ghrelin alone did not exert a significant effect on the expression of any of the genes or on NF- κ B luciferase activity, suggesting that like EM, ghrelin specifically reduced catabolic activities in response

to inflammatory stimuli. However, since EM, but not ghrelin, enhanced Col-II mRNA expression, but did not inhibit Col-X expression under IL-1 β treatment, it suggests that ghrelin has a different activity as compared to EM in terms of cartilage matrix gene expression. Using a serum-response element (SRE) reporter, which is an established reporter that serves as a readout of GHSR signaling as a G-protein coupled receptor [49, 50], we found that both ghrelin and EM indeed activated GHSR (Fig. 3C), suggesting that both ghrelin and EM are agonists at the ghrelin receptor.

Next, we determined whether ghrelin requires its receptor to inhibit IL-1 β activity by using primary chondrocytes from WT and GHSR null mice. We found that ghrelin inhibited IL-1 β -induced MMP13 expression in WT cells, but not in GHSR null cells (Fig. 3D). This result suggests that ghrelin, like EM, requires the ghrelin receptor to exert this function. We reasoned that if ghrelin and EM are both agonists at the ghrelin receptor, they would be competitors for the binding of this receptor. Thus, we treated mouse chondrocytes with EM and ghrelin simultaneously. Indeed, increasing concentrations of ghrelin led to a progressive decrease in the percentage of cells with EM presence in our immunocytochemistry analysis, thus further confirming that the ghrelin receptor is used by both EM and ghrelin (Fig. 3E). It is worth noting however, that the association of EM with chondrocytes could be overcome by a much lower concentration of ghrelin, suggesting that ghrelin is a much more potent competitor (Fig. 3E).

3.4. GHSR mediates the chondroprotective activity of EM *in vivo*

To determine whether GHSR is necessary to mediate the chondroprotective effect of EM *in vivo*, we subjected WT and GHSR null mice to intra-articular injection of monosodium iodoacetate (MIA). This model is well-established for recapitulating joint destruction and inflammation [51–56]. MIA is a specific inhibitor of glyceraldehyde-3-phosphate dehydrogenase (GAPDH), a key enzyme in the glycolysis pathway. Since articular cartilage is highly glycolytic as it adapts to its avascular environment [33, 34, 57], inhibition of glycolysis causes energy imbalance and cell death. As a result, there is a surge of inflammation. As MIA is cleared in the body, inflammation subsequently subsides to a large extent. However, as the case of injury-induced OA, already damaged cartilage continues to slowly disintegrate and enters the chronic phase of OA [51, 53–56]. To further validate the MIA model, we compared normal and human OA chondrocytes, and found OA chondrocytes to have reduced glucose uptake and ATP production as well as increased reactive oxygen species (ROS) production (Fig. 4A). These metabolic changes were also found in chondrocytes treated with MIA, suggesting MIA has caused oxidative stress as in human OA (Fig. 4B). Gene expression analysis indicated that MIA treatment increased pro-inflammatory cytokine IL-1 β and TNF α expression as well as hypertrophic chondrocyte marker Col-X and catabolic gene MMP13 expression in human articular chondrocytes (Fig. 4C). There were no differences in joint health between WT and GHSR null littermates under the control condition (PBS-injected), suggesting that the lack of GHSR does not affect the joint in the absence of OA (Fig. 4D). However, upon MIA treatment, GHSR null mice exhibited much more cartilage matrix loss than their WT littermates (Fig. 4D), suggesting that endogenous GHSR is required for preventing MIA-induced articular cartilage degeneration. We next determined whether GHSR mediates the chondro-protective activity

of EM *in vivo* by subjecting WT and GHSR null mice to intraperitoneal injection of EM, in addition to intra-articular injection of MIA. In WT mice, EM dramatically restored MIA-induced loss of safranin O staining, whereas EM could not restore matrix in GHSR null littermates (Fig. 4E). Similarly, EM could not rescue MIA-induced synovitis in the GHSR null joint as compared to WT (Fig. 4F). These data strongly demonstrate that GHSR mediates the chondroprotective activity of EM *in vivo*.

3.5. EM900, a macrolide devoid of antibiotic function, has a similar activity to EM and ghrelin

One concern of antibiotic use in disease treatment is antibiotic resistance. To determine whether the anti-inflammatory activity of EM is separate from the macrolide's antibiotic function, we used EM900 in *in vitro* and *in vivo* experiments. EM900 is a 12 membered EM derivative synthesized to be devoid of antibiotic activities, but has demonstrated anti-inflammatory activities in airway epithelial cells [58–60]. In our “live-dead” assay, chondrocytes showed good viability when treated with 1 and 10 μ M of EM900 (Fig. 5A), which were the concentrations subsequently used for the functional analysis of EM900. When EM900 was administered in pro-inflammatory cytokine IL-1 β -treated chondrocytes, a reduction of catabolic gene MMP13 expression was observed, suggesting that EM900 inhibited the catabolic activity of IL-1 β (Fig. 5B). Furthermore, the NF- κ B reporter activity induced by IL-1 β was also inhibited by EM900 (Fig. 5C), indicating that EM900 inhibits NF- κ B activation. To determine whether EM900 also activates the SRE luciferase reporter that can be activated by ghrelin and EM, we administered EM900 to human primary articular chondrocytes transfected with this reporter. Similar to ghrelin and EM, EM900 also led to the activation of SRE reporter (Fig. 5D).

To determine whether EM900 has a protective effect *in vivo*, we subjected normal mice to MIA-induced inflammatory osteoarthritis, and intraperitoneally administered EM900. Histological analyses showed significant level of cartilage matrix loss in MIA-treated knee joints, but much less matrix loss when EM900 was also injected, which was subsequently quantified using the OARSI scoring system (Fig. 5E). Consistently, EM900 also significantly reduced MIA-induced synovitis induced by MIA (Fig. 5F). While it is not clear whether the ghrelin receptor is required to mediate the activity of EM900, our results demonstrate that EM900 has an anti-inflammatory and joint protective activity similar to EM in isolated human chondrocytes as well as in an inflammatory OA animal model, and suggests that this activity is antibiotic-independent.

3.6. EM, EM900 and ghrelin enhance cartilage matrix levels in human OA cartilage

Although EM, EM900 and ghrelin have demonstrated inhibitory activities of IL-1 β in the *in vitro* cultures of primary human chondrocytes, whether they could promote cartilage matrix levels in human cartilage under diseased conditions was not known. Thus, we subjected OA cartilage specimens to EM and ghrelin treatment. As expected, these OA cartilage samples also showed reduced expression of cartilage matrix genes aggrecan and collagen IX and had increased expression of catabolic genes MMPs and IL-1 β (Fig. 6A). Significantly, ghrelin expression was reduced in OA cartilage specimens (Fig. 6A). When administered in *ex vivo*-cultured OA articular cartilage explants, EM, EM900 and ghrelin all led to significantly

increased safranin O staining as compared to PBS control samples (Fig. 5B–5D), suggesting that EM, EM900, as well as ghrelin indeed enhance cartilage matrix levels even in advanced OA.

4. Discussion

This study provides a key mechanism for the pharmacological activity of erythromycin (EM). We report that the ghrelin receptor is an essential mediator of EM chondroprotective activity, demonstrating that both ghrelin and EM can activate the ghrelin receptor and inhibit pro-inflammatory cytokine IL-1 β -induced NF- κ B activation and catabolic gene expression. Furthermore, the ghrelin receptor is required to resist synovitis and joint destruction in EM-treated mice *in vivo*, thus playing an essential role for joint maintenance under pathological conditions.

Beneficial effects of ghrelin on OA cartilage has only recently been reported. In 2017, Zou *et al.* found that the synovial fluid of knee OA patients contained lower levels of ghrelin [61]. In the same year, Qu *et al.* showed that ectopic intraperitoneal injection of ghrelin could reduce surgery-induced mouse knee cartilage damage and inflammation [31]. In 2018, Liu *et al.* also demonstrated that supplementing ghrelin in culture media inhibited IL-1 β -induced catabolic changes in human cartilage [30]. Our study is consistent with these recent findings, but we have also significantly deepened the understanding on the role of ghrelin in OA in additional ways. First, we definitely demonstrate that ghrelin signaling is *critical* for joint health and resistance to inflammation. Past reports focused on ectopic ghrelin treatment and did not indicate whether ghrelin signaling is required for the maintenance of the joint [30, 31]. Our study demonstrated that ghrelin signaling is essential for resisting inflammatory cytokine-induced MMP expression *in vitro* and delaying cartilage destruction *in vivo*. Second, we showed that ghrelin enhances cartilage matrix levels in human OA cartilage, even in the absence of exogenous pro-inflammatory cytokine. This has important implications for the potential use of ghrelin or its agonists to treat existing OA and halt cartilage destruction. Third, we extended our analysis *in vivo* to the analysis of the synovium, which plays an important role in inflammation of the joint, rather than being restricted to cartilage analysis. It is worth noting that while ghrelin administration would increase feeding and body weight [62, 63], mice lacking ghrelin have the same weight as the wild type under normal feeding conditions [64, 65]. On the other hand, mice lacking the receptor for ghrelin showed sexual dimorphism in this aspect: male mice did not exhibit any difference in body weight differences when fed with the standard chow, but female mice did [26]. Since we only used male mice, it allays the concern of body weight-associated OA in our study.

One important discovery of our study is the identification of a pharmacological mechanism for the macrolide antibiotic erythromycin (EM) in promoting joint health. EM is a commonly used antibiotic that directly binds bacterial ribosomes to inhibit protein synthesis [66]. Although EM-like macrolide antibiotics have anti-inflammatory activities in the lung and bone in an antibiotic-independent manner, its chondroprotective effect in the joint has only been described in our recent study [17–22]. However, in all these systems, how EM exerts its effect on eukaryotic cells had not been elucidated. EM was previously shown to be

capable of binding to the motilin receptor [67, 68]. The natural ligand of the motilin receptor is motilin, a peptide that stimulates GI motility [69]. Interestingly, in the mouse, motilin and its receptors are pseudogenes [70, 71]. Since our *in vivo* experiments indicate a joint protective effect of EM, we reasoned that the motilin receptor could not be mediating the activity of EM in the mouse [22]. The ghrelin receptor and the motilin receptor share 50% identity, and both ligands have very similar roles in promoting gastric emptying and GI motility [69, 72, 73]. While an EM derivative binds to the ghrelin receptor, it is still not known whether EM itself can directly bind to the receptor. Nevertheless, our subsequent experiments indicated that ghrelin can compete with EM in ghrelin receptor association, and that the effect of EM is lost when the ghrelin receptor is not present. Thus, our data strongly suggest that EM functionally acts through the ghrelin receptor to exert its activity in the mouse. However, this does not exclude the motilin receptor as a mediator of the activity of EM in species where motilin signaling is present, as in humans [70]. While the binding affinity of EM to the ghrelin receptor *in vivo* and *in vitro* is now known, our data indicate that the association of EM with chondrocytes can be outcompeted by a much lower concentration of ghrelin, suggesting that it is not likely that EM will inhibit the activity of endogenous ghrelin (Fig. 3F). Furthermore, in the case of OA, when ghrelin concentration is already lower in the synovial fluid [61], additional EM might stimulate the ghrelin receptor to exert its beneficial effect. However, in order to select the most appropriate macrolide for OA therapy, other EM-like antibiotics, such as azithromycin or clarithromycin, would also need to be investigated for similar effects on the joint as EM,

Systemic administration of macrolide antibiotics may raise the concern of antibiotic resistance. However, this concern may be allayed by intra-articular injection of antibiotics or by administration of non-antibiotic macrolides such as EM900. One limitation of this study is that we have not demonstrated whether EM900 acts through the ghrelin receptor to exert its activity, even though we have shown this is the case for EM. Another limitation is that only one mouse model for joint inflammation and destruction was tested. Since there are many causes of OA, it will be important to evaluate the role of these reagents in other models, such as the Destabilization of the Medial Meniscus (DMM), meniscectomy, or obesity-induced OA model [74–78].

Future studies will also involve the delivering of EM into OA joints after OA has initiated for a period of time so that its therapeutic potential can be evaluated. The only study on EM in human OA was restricted to a single clinical trial combining EM with acetaminophen [79]. While joint effusion and pain was reduced [79], changes in joint structure were not assessed and erythromycin efficacy as a monotherapy was also not determined [79]. Our study has characterized the structural changes of the joint and uncovered a new mechanism for EM to act as an anti-inflammatory reagent. While ghrelin has been reported to alleviate pain in the carrageenan intraplantar injection model [80], it remains unclear whether such parallels exist in the knee joint. Thus, future experiments would be designed to determine whether EM and ghrelin can provide pain relief in arthritic joints, to thoroughly explore the potential of these molecules as OA therapeutics.

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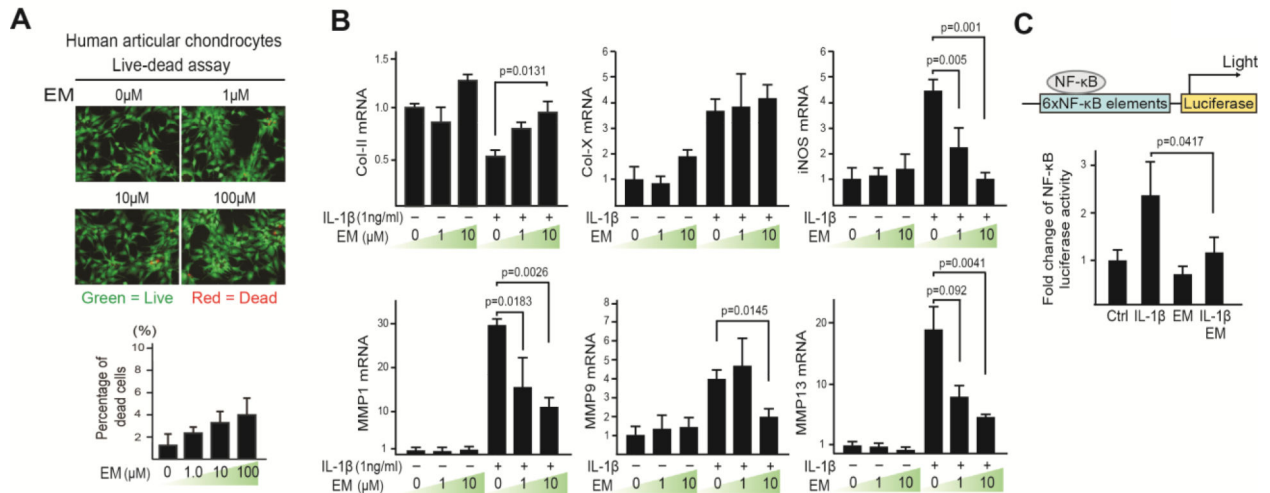


Fig. 1. Erythromycin (EM) inhibits the catabolic activity of IL-1 β on primary human articular chondrocytes (nHACs).

(A) Live-dead assay on chondrocytes treated with EM (1, 10 and 100 μ M) for 4 days.

Percentage of dead cells/total cells was quantified. (B) RT-qPCR analysis of Col-II, Col-X, iNOS, MMP1, MMP9 and MMP13 on nHACs treated with EM (0, 1, and 10 μ M) and IL-1 β (1ng/ml). TBP served as a reference gene for normalization. (C) NF- κ B transactivation assay. nHACs were transiently transfected with an NF- κ B reporter and treated with EM

(10 μ M) and IL-1 β (1ng/ml). A Renilla luciferase construct was co-transfected as an internal control for normalization. Data were reported as mean \pm SD and analyzed by Dunnett's test (B) and an unpaired *t*-test (C). At least three independent experiments were performed. *

$p < 0.05$.

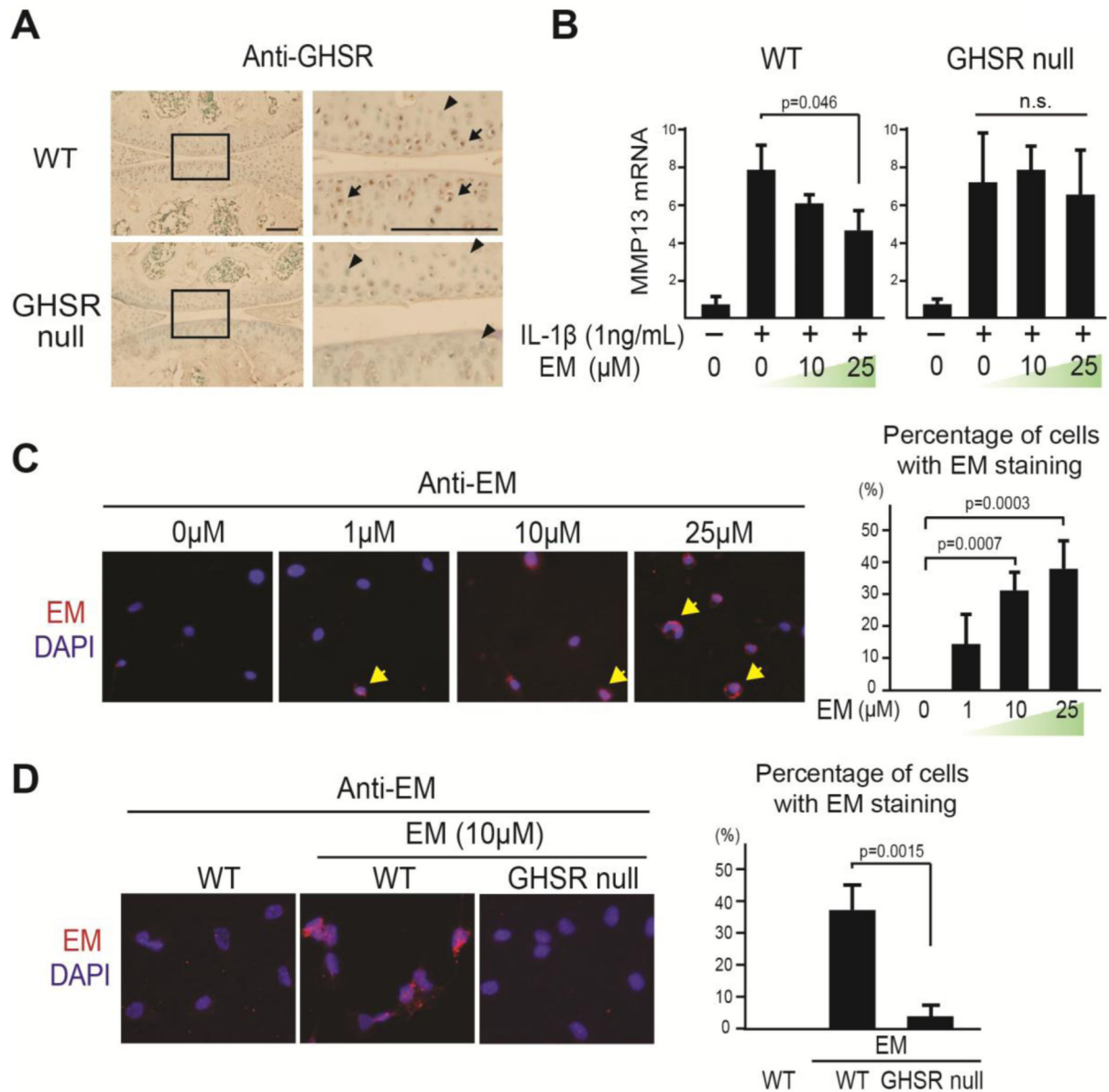


Fig. 2. The anti-inflammatory activity of EM in chondrocytes is mediated by the ghrelin receptor GHSR.

(A) Immunohistochemical (IHC) analysis using an anti-GHSR antibody (anti-GHS-R1a) on WT and GHSR null mouse knee joints. Rectangles denote areas magnified. Arrows and arrowhead indicate GHSR positive and negative cells, respectively. Scale bar = 200 μ m. (B) Relative MMP13 mRNA analysis on WT and GHSR null chondrocytes treated with EM (0, 10, and 25 μ M) and IL-1 β (1ng/ml). TBP served as a loading control. (C) Immunocytochemistry (ICC) analysis with anti-EM antibody (red) on WT chondrocytes treated with different doses of EM (0, 1, 10, and 25 μ M) for 1hr. Arrows indicate cells with EM staining. Percentage of EM-positive cells was calculated. (D) ICC analysis with EM presence on WT and GHSR null chondrocytes treated with EM (10 μ M) for 1hr. Percentage of EM-positive cells was calculated. Data were reported as mean \pm SD and analyzed by

Dunnett's test (B and C) and an unpaired *t*-test (D). At least three independent experiments were performed. * $p < 0.05$. n.s: not significant.

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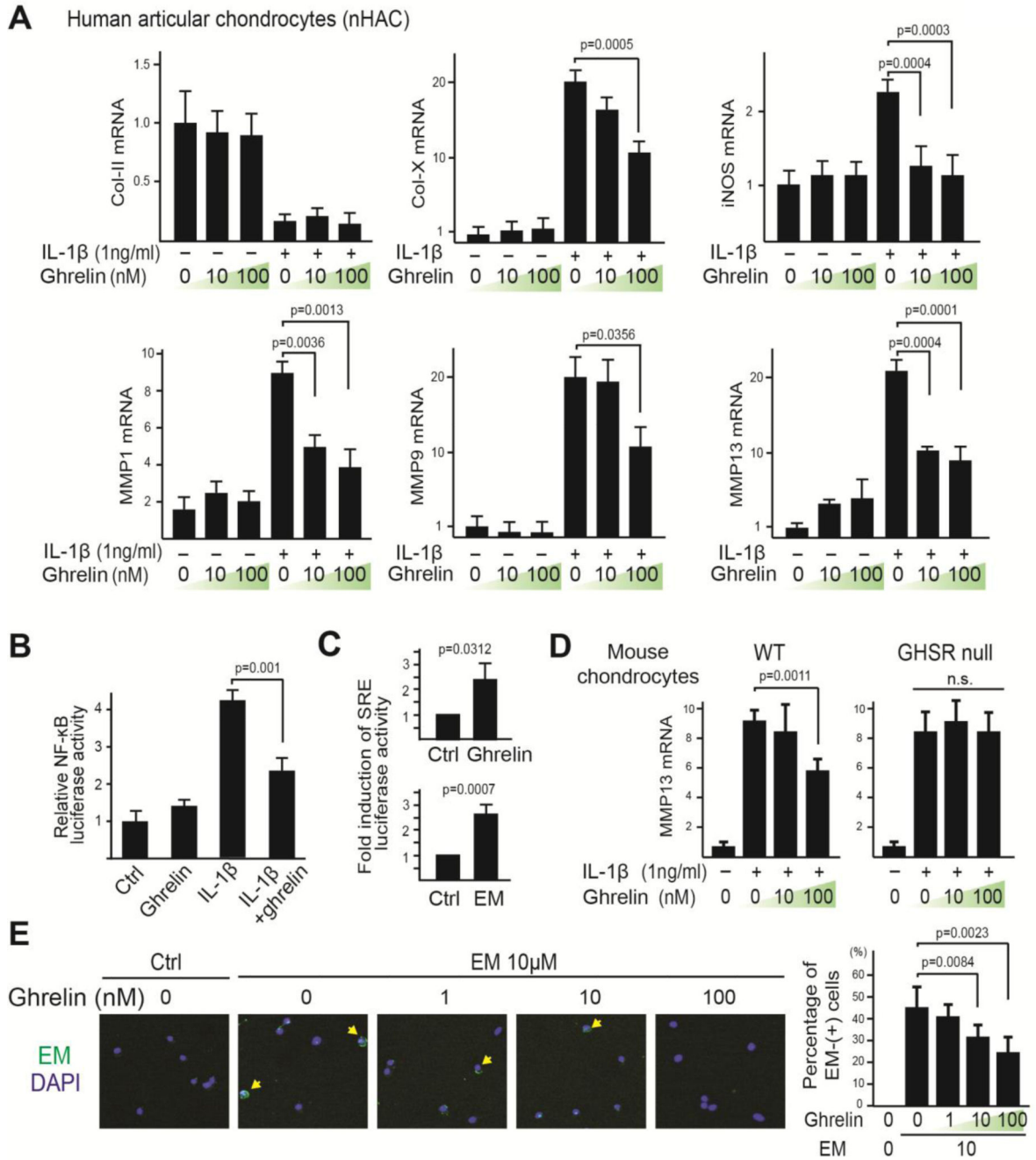


Fig. 3. Ghrelin signaling is necessary and sufficient to inhibit IL-1 β -induced catabolic gene expression and NF- κ B activation in chondrocytes.

(A) RT-qPCR analysis of Col-II, Col-X, iNOS, MMP1, MMP9 and MMP13 on nHACs treated with ghrelin (0, 10 and 100nM) and IL-1 β (1ng/mL). TBP served as a reference gene. (B) NF- κ B transactivation assay. nHACs were transiently transfected with NF- κ B reporter construct for 24 hours and then treated with ghrelin (10nM) and IL-1 β (1ng/mL). A Renilla luciferase construct was co-transfected as an internal control for normalization. (C) Serum Response Element (SRE) luciferase reporter assay. nHACs were transiently transfected with the SRE reporter construct for 24 hours and then treated with ghrelin

(10nM) or EM (10 μ M) for 16 hours. A Renilla luciferase construct was co-transfected as an internal control for normalization. Data are presented as fold activation relative to untreated samples. (D) RT-qPCR analysis of MMP13 on WT and GHSR null mouse chondrocytes treated with ghrelin (0, 10, 100nM) and IL-1 β (1ng/mL). TBP served as a reference gene. (E) ICC analysis with an anti-EM antibody on WT chondrocytes co-treated with EM (10 μ M) and different concentrations of ghrelin (1, 10 and 100nM). Arrows indicate cells with EM staining. Percentage of EM-positive cells was calculated. Data were reported as mean \pm SD and analyzed by Dunnett's test (A, D and E) and an unpaired *t*-test (B and C). At least three independent experiments were performed. * *p*<0.05. n.s: not significant.

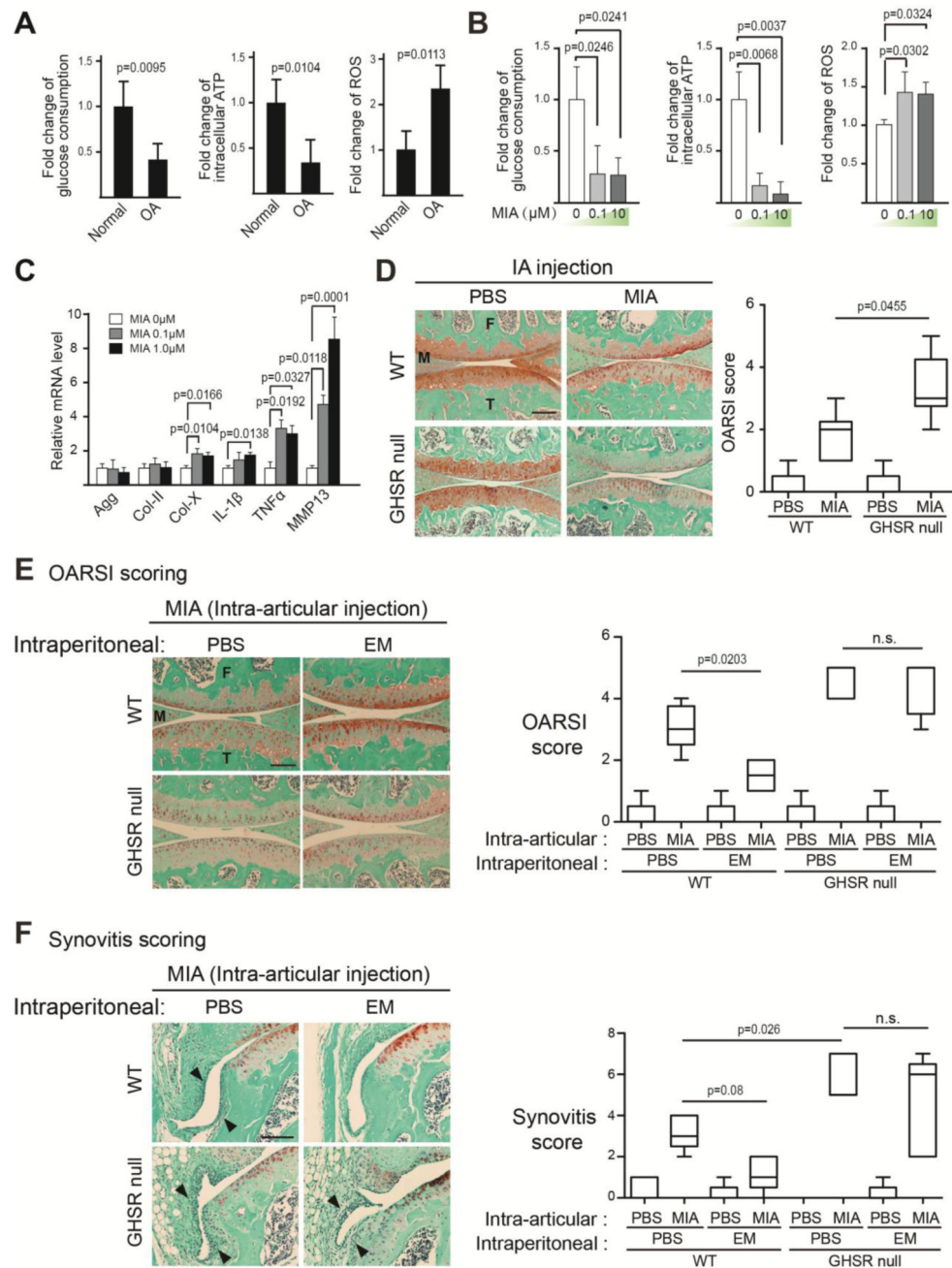


Fig. 4. The Ghrelin receptor is required for EM to inhibit cartilage matrix loss and synovitis *in vivo*.

(A) Analysis of total glucose consumption, intracellular ATP levels and reactive oxygen species (ROS) levels at day 2 of culture in healthy and OA human articular chondrocytes. (B) Analysis of total glucose consumption, intracellular ATP levels and reactive oxygen species (ROS) levels at day 2 of culture in nHACs after MIA treatment. (C) RT-qPCR analysis of aggrecan (agg), Col-II, Col-X, IL-1 β , TNF α and MMP13 on nHACs treated with MIA (0, 0.1 and 1 μ M) for 2 days. TBP served as a reference gene for normalization. (D)

Safranin O staining analysis of WT and GHSR null mouse knee joints. MIA (50µg/joint) was intra-articularly injected to knee joints at day 0, and samples were harvested at day 7. The degree of cartilage matrix loss was examined according to the OARSI scoring system. **(E)** Safranin O staining analysis of WT and GHSR null mouse knee joints injected with MIA. MIA was intra-articularly injected to knee joints at day 0. EM (PBS as control) was IP-injected at 50µg/g into the mice daily until sample harvest at day 7. The degree of cartilage matrix loss was examined according to the OARSI scoring system. **(F)** Synovitis analysis on sections from the medial tibial plateau of WT and GHSR null mice knees injected with EM. MIA was intra-articularly injected to knee joints at day 0. EM (PBS as control) was IP-injected into the mice daily until sample harvest at day 7. Arrowheads indicate thickening of the synovial layers and increased cellularity of resident cells. M = Meniscus. T = Tibia. F=Femur. Synovitis was scored according to established synovitis scoring protocols. Scale bar = 200µm. For A, data were reported as mean ± SD and analyzed by an unpaired *t*-test. For B and C, data were reported as mean ± SD and analyzed by Dunnett's test. Comparisons were made between MIA treatments over untreated controls for each gene. For D-F, data were reported as a box plot, where the median, as well as the maximum and the minimum data points were indicated. Data from D-F were analyzed by Kruskal-Wallis statistical test followed by Mann-Whitney *U* test. * $p < 0.05$. n.s: not significant.

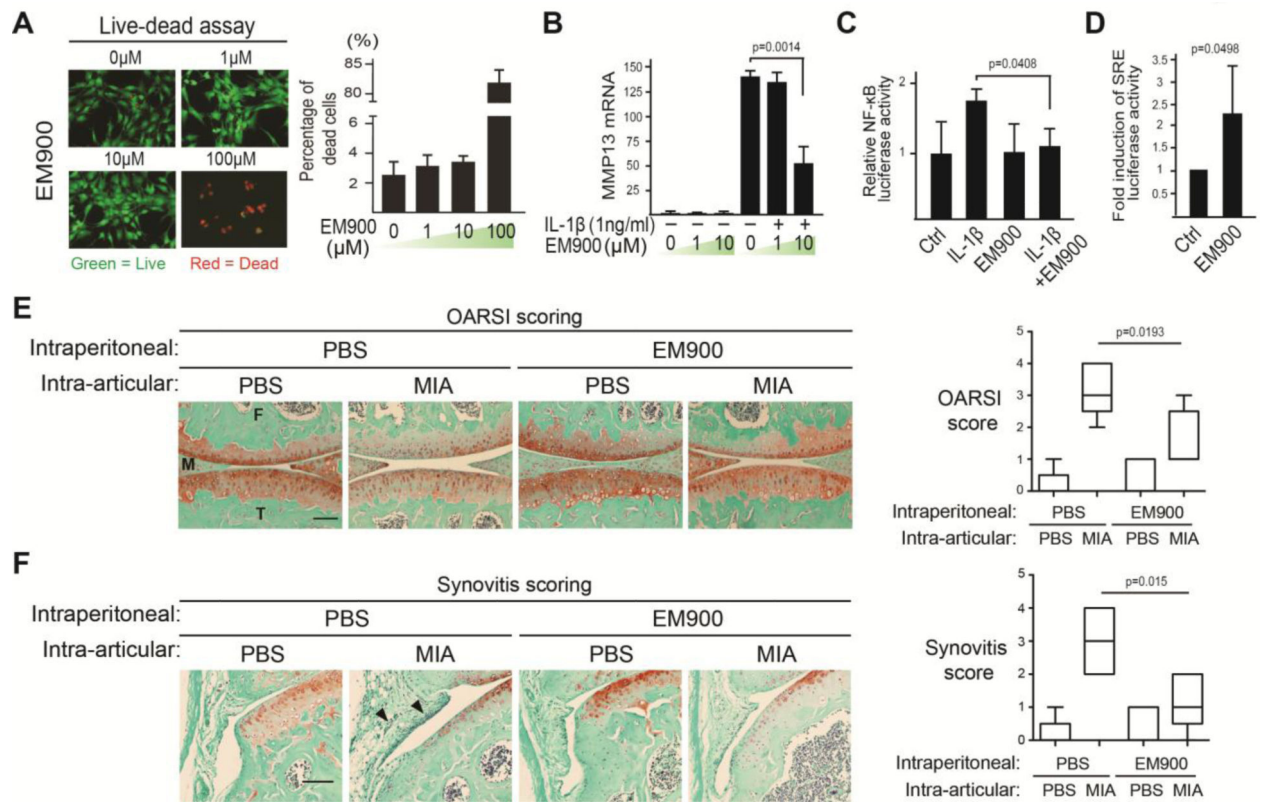


Fig. 5. EM900, a non-antibiotic EM derivative, inhibits IL-1 β -induced MMP13 expression and NF- κ B activation, and maintains joint health in MIA-injected knees.

(A) Live-dead assay on chondrocytes treated with EM900 (0, 1, 10 and 100 μ M) for 4 days. Percentage of dead cells/total cells was quantified. (B) RT-qPCR analysis of MMP13 mRNA on nHACs treated with EM900 (1, 10 and 25 μ M) overnight followed by IL-1 β (1ng/mL) treatment for 3 days. TBP served as a reference gene. (C) NF- κ B transactivation assay. nHACs were transiently transfected with the NF- κ B reporter construct for 24 hours and then treated with EM900 (1 μ M) and IL-1 β (1ng/mL). A Renilla luciferase construct was co-transfected as an internal control for normalization. (D) Serum Response Element (SRE) luciferase reporter assay as a readout of ghrelin receptor signaling. nHACs were transiently transfected with the SRE reporter construct for 24 hours and then treated with EM900 (10 μ M) and IL-1 β (1ng/mL). A Renilla luciferase construct was co-transfected as an internal control for normalization. Data were presented as “fold activation” compared to untreated samples. (E) Safranin O staining images for cartilage matrix analysis of WT mouse knee joints. MIA was intra-articularly injected to knee joints at day 0. EM900 was IP-delivered daily from the day of MIA injection. Samples were collected for analysis 7 days later. The degree of cartilage matrix loss was scored according to the OARS1 scoring system. Scale bar = 200 μ m. M = Meniscus. T = Tibia. F = Femur. (F) Synovitis analysis on sections from the medial tibial plateau of the WT mice knees injected with EM900. MIA was intra-articularly injected into knee joints at day 0. EM900 (PBS as control) was IP-injected into the mice daily until sample harvest at day 7. Arrowheads indicate thickening of the synovial layers and increased cellularity of resident cells. Synovitis was scored according to established synovitis scoring protocols. For A to D, data were reported as mean \pm SD and analyzed by

Dunnett's test (B) and an unpaired *t*-test (C, D). For E and F, data were reported in a box plot, where the median, as well as the maximum and the minimum data points were indicated. Data from E and F were analyzed by Kruskal-Wallis statistical test followed by Mann-Whitney *U* test. * $p < 0.05$.

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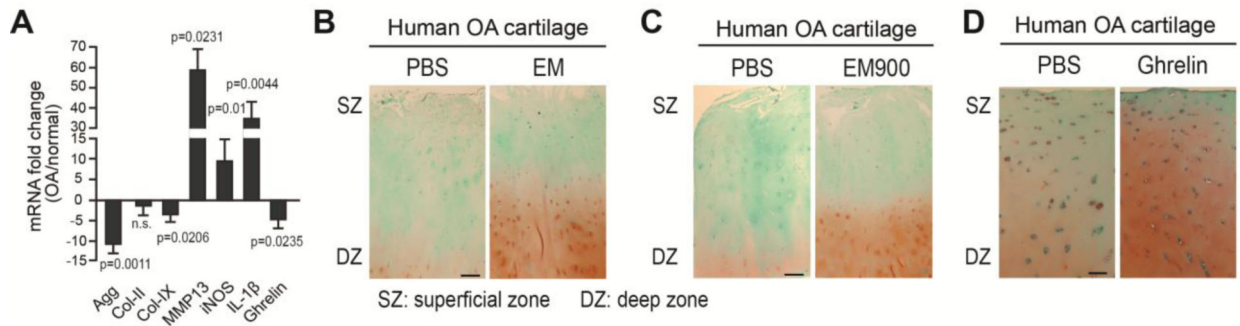


Fig. 6. EM and ghrelin enhance cartilage matrix levels in human OA cartilage.

(A) RT-qPCR analysis of Agg, Col-II, Col-IX, MMP13, iNOS, IL-1 β and ghrelin in human normal cartilage from 4 healthy donors and OA cartilage from 5 OA donors. TBP served as a reference gene. (B) Safranin O staining analysis on sections of OA articular cartilage explants after 3 weeks of culturing with 10 μ M EM. Images of explants from donor 54/F (age/sex) are shown. (C) Safranin O staining analysis on sections of OA articular cartilage explants after 3 weeks of culturing with 10 μ M EM900. Images of explants from donor 54/F (age/sex) are shown. (D) Safranin O staining analysis on sections of OA articular cartilage explants after 3 weeks of culturing with 100nM ghrelin. Images of explants from donor 53/F (age/sex) are shown. For each experiment, duplicate technical repeats were analyzed. Experiments were repeated three times using samples from three different OA donors. As the extent of cartilage matrix loss and OA severity of the donor samples differed among experiments, cellularity and matrix levels varied, which is evident from the differences in control samples used for EM, EM900 and ghrelin experiments. Scale bar = 100 μ m. SZ = superficial zone. DZ = deep zone.