**Conditional deletion of E11/podoplanin in bone protects against load-induced osteoarthritis**

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**Abstract**

**Background:** Subchondral bone (SCB) thickening is one of the earliest detectable changes in osteoarthritic joints and is considered a potential trigger for subsequent articular cartilage degeneration. In this manuscript, we examine whether disruption to the SCB osteocyte network contributes to the initiation and pathogenesis of osteoarthritis.

**Methods:** We examined expression patterns of the glycoprotein E11/podoplanin by immunohistochemical labelling in murine, human and canine osteoarthritis models. We also examined the effects of twice-weekly administration of Bortezomib, a proteasome inhibitor which stabilises osteocyte E11 levels, to C57/BL6 wild-type male mice (1mg/kg/day) for 8 weeks after surgical destabilisation of the medial meniscus. By inducing osteoarthritis-like changes in the right knee joint of 12‐week‐old male E11 hypomorphic mice (and corresponding controls) using a post-traumatic joint loading model, we also investigated whether a bone-specific E11 deletion in mice increases joint vulnerability to osteoarthritis. Articular cartilage degradation and osteophyte formation were assessed by histology and in line with the OARSI grading system.

**Results:** Our studies reveal increased E11 expression in osteocytes of human and canine osteoarthritic SCB. We found that Bortezomib administration had no effect on surgically-induced osteoarthritis, potentially due to a lack of the expected stabilisation of E11 in the SCB. We also found, in concordance with our previous work, wild-type mice exhibited significant load-induced articular cartilage lesions on the lateral femoral condyle (p<0.01) and osteophyte formation. In contrast, E11 hypomorphic mice did not develop osteophytes or any corresponding articular lesions.

**Conclusions:** Overall, these data suggest that an intact osteocyte network in the SCB contributes to the development of mechanically-driven osteoarthritis. Further, the data presented here indicate that the molecular pathways that preserve the osteocyte network, such as those driven by E11, may be targeted to limit osteoarthritis pathogenesis.

**Key words:** osteocytes,subchondral bone, osteoarthritis, E11/podoplanin

**Background**

Osteoarthritis is a degenerative joint disease and a global health care burden. In osteoarthritis, the articular cartilage undergoes structural deterioration, causing joint pain, loss of joint function and significantly reducing quality of life. However, its underlying molecular mechanisms are not fully understood. As such, there is an ever-growing need for an effective disease-modifying treatment.

Although often considered secondary, subchondral bone (SCB) thickening in osteoarthritic joints is one of the earliest detectable changes and is now considered a potential trigger for subsequent articular cartilage degeneration [1, 2]. Osteoblast-derived osteocytes are the most numerous of all the cells within bone and have a unique morphology with extensive dendritic processes creating bone’s osteocyte-canalicular network. This network is now known to orchestrate bone remodelling [3]. However, in osteoarthritic joints, the osteocytes in the SCB exhibit alterations to their exquisite dendritic morphology, with fewer and more disorganised dendrites [4]. Furthermore, other reports have noted that the expression of sclerostin, the mature osteocyte marker, is disrupted in osteoarthritic SCB [5, 6]. Together, these data suggest that the osteocyte may contribute a central role to pathological SCB sclerosis in osteoarthritis and that an intact osteocytic network is necessary for maintaining healthy SCB architecture.

Numerous genes have been suggested to influence osteocyte formation, one of which encodes the transmembrane glycoprotein E11/podoplanin. We and others have previously shown that E11 is expressed by early embedding osteocytes, thus identifying it as a factor which likely contributes to the vital, early stages of osteocyte differentiation [7–9]. It is known that mechanical strain *in vivo* increases E11 mRNA expression [7] and that E11 siRNA abrogates the formation of osteocyte dendrites. [7]. In contrast, over-expression of E11 in ROS 17/2.6 osteoblast-like cells has been found to promote the formation of long dendritic processes [10–12]. Furthermore, we have recently reported that E11 levels are regulated post-translationally by proteasomal degradation and that their preservation, through the administration of proteasome inhibitors such as Bortezomib, leads to the induction of an osteocyte-like morphology in MLO-A5 pre-osteocytic cells [9]. In accordance with this, we recently showed that the hypomorphic bone-specific ablation of E11 in mice results in disrupted osteocytic dendrite formation, which supports a key role for E11 in regulating the cytoskeletal changes associated with osteocyte process formation and elongation [13].

As the formation of such dendritic processes is a key functional feature of the normal mature osteocyte network, which is perturbed in osteoarthritis [4], we have examined herein whether disruption to the integrity of the osteocyte dendritic processes contributes to the initiation of osteoarthritis. Specifically, we investigated whether proteasome inhibition can stabilise E11 expression *in vivo* to protect against the osteoarthritis that develops following surgically-induced destabilisation of the medial meniscus (DMM). Moreover, we examined whether the bone-specific conditional deletion of E11 in mice affects early adaptive processes and joint vulnerability to osteoarthritis induction by a mechanically-induced post-traumatic osteoarthritis model.

**Methods**

*Animals*

C57/BL6 mice harbouring a conditional deletion of E11 in late osteoblasts (osteocalcin promoter driven; cKO; hypomorphic with ~70% reduction in E11 protein expression) as well as their appropriate *E11*flox/flox control littermates (WT) were kept in polypropylene cages, with light/dark 12-h cycles, at 21 ± 2°C, and fed *ad libitum* with maintenance diet (Special Diet Services, Witham, UK) [13]. We obtained floxed Pdpn mice from the UCOMM/KOMP, MRC Harwell, Oxfordshire, UK and osteocalcin-cre mice as a kind gift from Thomas Clemens at John Hopkins Medicine, Baltimore, Maryland. cKO and WT mice were generated as described previously [13]. All analyses were conducted blindly to minimise the effects of subjective bias. All experimental protocols were approved by Roslin Institute's Animal Users Committee and the animals were maintained in accordance with UK Home Office guidelines for the care and use of laboratory animals.

*In vivo joint loading*

12-week old male cKO (n=5) and WT (n=3) mice were isoflurane-anaesthetised (4% and maintained at 2% during loading) and the right knee joint loaded as described previously [14]. Briefly, using a servo-electric materials testing machine (Electroforce 3100, Bose, UK), axial compressive loads were applied through the right knee joint via custom-made cups. All studies used a single loading pattern in which peak loads of 11N for the cKO and 12N for the WT mice (see [13]) were applied (for 0.05 seconds; 0.025 seconds rise and fall time; 9.9 seconds baseline hold time at between periods of peak loading). Joints were loaded for 40 cycles, 3 times/week for 2 weeks in the morning and the left (non-loaded control) and right knees dissected 3 days after the final loading episode. Knee joints were fixed in 4% paraformaldehyde for 24 hours at 4oC before being stored in 70% ethanol. Mice were sacrificed by exsanguination and confirmation of death by cervical dislocation.

*Destabilisation of the medial meniscus (DMM)*

Osteoarthritis was induced in 8-week old C57/BL6 male mice (Charles River) by surgically induced DMM under isoflurane-induced anaesthesia (see above). Animals were randomly allocated to treatment groups to reduce subjective bias. We elected not to perform sham surgery on the contralateral knee based on animal welfare grounds since previous studies had shown no difference in osteoarthritis scores between non-operated and sham-operated knee joints using this model and since the primary aim of the experimental research was to evaluate the potential protective effect of Bortezomib following DMM [15–17]. Following transection of the medial meniscotibial ligament, the joint capsule and skin were closed and anaesthesia reversed. Mice then either received twice-weekly morning intraperitoneal injections of Bortezomib (1mg/kg [18]; n=8) or vehicle control (99.7% w/v saline; n=8) for 8 weeks at which point knee joints were dissected, fixed in 4% paraformaldehyde for 24 hours at 4oC, and then stored in 70% ethanol. Mice were sacrificed by exsanguination and confirmation of death by cervical dislocation.

*Micro-computed tomography (microCT) analysis*

Scans were performed with an 1172 X-Ray microtomograph (Skyscan, Belgium) to evaluate the SCB. High-resolution scans with an isotropic voxel size of 5 µm were acquired (50 kV, 200µA, 0.5 mm aluminium filter, 0.6° rotation angle). The projection images were reconstructed using NRecon software version 1.6.9.4 (Skyscan, Belgium). Each dataset was rotated in Dataviewer (Skysan, Belgium) to ensure similar orientation and alignment for analysis. Hand-drawn regions of interests (ROI) of the SCB trabecular bone for each femur/tibia lateral/medial compartments was first achieved [19]. SCB ROIs was subsequently selected for each compartment. Analysis of SCB plate thickness and the epiphyseal trabecular bone was achieved using 3D algorithms in CTAn (Skyscan, Belgium) to provide: SCB plate thickness (SCB Th.; mm); epiphyseal trabecular bone volume/tissue volume (Tb. BV/TV; %); trabecular number (Tb. N.; mm-1); trabecular thickness (Tb. Th.; mm); trabecular separation (Tb. Sp.; mm); trabecular pattern factor (Tb. Pf.: mm-1) .

*Human and animal osteoarthritic samples*

The human SCB samples were obtained from patients undergoing total knee replacement for osteoarthritis. Samples (discarded femoral and tibial bone cuts) were obtained with patient consent and ethical approval from the NHS Lothian Bioresource. The collection, storage, and subsequent use of human tissues are regulated in Scotland by The Human Tissue Act (Scotland) 2006. Canine osteoarthritis samples were residual tissues collected from pets undergoing surgery for the treatment of elbow joint disease with informed consent (osteoarthritis), or which had died of unrelated disease (healthy). Consent for use was obtained from the animal owners and ethical approval for their collection and use given by the Veterinary Ethical Review Committee of the University of Edinburgh (VERC; approval 23/12). In all cases, the joints were macroscopically evaluated for signs of osteoarthritis.

*Histological analysis*

Murine left and right knee joints were decalcified, wax-embedded and 7μm coronal sections cut. For assessment of osteoarthritis severity, multiple sections (five/slide) from 120 μm intervals across the whole joint were stained with Toluidine blue (0.4% in 0.1 M acetate buffer, pH 4) and counterstained with Fast Green (0.2% in dH2O). Articular cartilage lesion severity was graded using the well-established OARSI grading scale [20]. Scoring was conducted blindly with a second observer scoring ~25% of the sections. For assessment of osteoclast activity, slides were stained with Goldner’s Trichrome using standard procedures or for tartrate resistant acid phosphatase (TRAP). For TRAP staining 70mg napthol AS-TR phosphate (Sigma) was dissolved in 250µl N-N dimethyl formamide (Sigma) and added to 50ml 0.2M sodium acetate buffer pH5.2. 115mg sodium tartrate dihydrate (Sigma) and 70mg fast red salt TR (Sigma) was dissolved into this solution and slides were incubated at 37oC for 2 hours. Sections were counterstained in Meyer’s haematoxylin (Sigma), washed in distilled water and mounted in aqueous mounting medium (Vector Labs). Slides were imaged using a NanoZoomer slide scanning system (Hamamatsu) and histomorphometry performed using Bioquant Osteo (Bioquant Image Analysis Corporation).

*Immunohistochemistry*

For immunohistochemical localisation of E11 and sclerostin, sections were dewaxed in xylene and rehydrated. Sections were incubated at 37 °C for 30 min in 1mg/ml trypsin for antigen demasking. Endogenous peroxidases were blocked by treatment with 3% H2O2 in methanol (Sigma). Species-specific amino acid sequences for E11 and sclerostin were obtained and Clustal Omega was used to perform alignment and comparison of the mouse, human and dog sequences. The level of homology between amino acid sequences for two different species was expressed as mean percent identity (Suppl. Table. 1). As a result of this and following optimisation, antibodies used were: E11 mouse samples (IgG polyclonal raised in goat; R&D systems; 1/100); E11 human and canine samples (IgG polyclonal raised in sheep; R&D systems; 1/500). Sclerostin mouse samples (IgG polyclonals raised in goat; R&D systems; 1/200); with appropriate controls [9]. The Vectastain ABC universal detection kit (Vector Laboratories, Peterborough, UK) was used according to the manufacturer’s instructions. The sections were finally dehydrated, counterstained with haematoxylin and mounted in DePeX. All sections to be compared were immunostained at the same time in order to standardise conditions and minimise any differences in antibody incubation times. Positive E11 and sclerostin staining in the articular cartilage chondrocytes and SCB osteocytes was readily identifiable and we were therefore able to semi-quantify staining intensity in these cell types between the different treatment groups.

*Statistical analysis*

Statistical analysis of articular cartilage lesion grades compared loaded/DMM (right) and contra-lateral control (left) joints by paired Wilcoxon's signed-rank test. P < 0.05 was considered statistically significant.

**Results**

*E11 expression is increased in human and canine osteoarthritic SCB osteocytes*

We first sought to examine the expression of E11 in the SCB of two different osteoarthritis animal models, and in samples from patients undergoing total knee replacement (Fig. 1). Immunohistochemical labelling revealed that E11 expression was similar in the SCB of both surgically-induced DMM and non-operated mice in both the lateral and medial aspects of the joint (arrows, Fig. 1A). E11 immunolabelling was also observed in the articular cartilage chondrocytes (Fig. 1A). However, increased E11 expression was observed in the SCB osteocytes in naturally occurring human (Fig. 1B) and canine (Fig. 1C) osteoarthritis in comparison to unaffected control tissue. This indicates that E11 expression levels are raised in osteoarthritic SCB osteocytes, suggesting that this may be linked to pathology.

*Administration of Bortezomib has no effect on surgically induced osteoarthritis*

We have previously shown that exposure to the proteasome inhibitor, Bortezomib, stabilised E11 expression *in vitro* [9]*.* Based upon this observation, we hypothesised that Bortezomib treatment would protect against osteoarthritis pathology *in vivo* through an enhanced stabilisation of E11 and the promotion of osteocyte differentiation. We found no significant differences in the weights of mice treated with Bortezomib in comparison to vehicle-treated mice (Fig. 2A). There was also no significant difference in either the maximum (Fig. 2B) or mean (Fig. 2D) OARSI osteoarthritis scores for the non-operated joints between vehicle and Bortezomib treated mice. The DMM-operated joints showed an expected increase in the OARSI scores in comparison to the non-operated joints (compare Fig. 2C & E to 2B & D) however, there were no significant differences in the OARSI scores between vehicle-treated and Bortezomib-treated joints with DMM (Fig. 2C, E & F).

In order to determine if there were any SCB abnormalities after DMM surgery in Bortezomib treated mice, we performed microCT analysis. No significant DMM-related differences were observed in the medial tibia SCB plate and epiphyseal trabecular bone parameters (Fig. 3A – F). In contrast, Bortezomib treated mice exhibited significant DMM-related increases in the SCB thickness (P<0.05, Fig. 3G) and trabecular thickness (P<0.05, Fig. 3J) in the lateral tibia, in comparison to vehicle-treated mice. No significant differences were observed in the other lateral tibia epiphyseal parameters.

To assess whether Bortezomib did indeed stabilise E11 expression, we performed immunolabelling for E11 in both non-operated control and DMM-operated knee joints with or without Bortezomib treatment. In the vehicle-treated joints, E11 was expressed in the SCB osteocytes as expected, and also in the chondrocytes of the superficial and middle zones of the articular cartilage (Fig. 4A). In accordance with our results in Fig. 1A, DMM had no effect on the SCB expression of E11 (Fig. 4A). The administration of Bortezomib, however, provoked decreases in the articular cartilage chondrocyte expression of E11, with no apparent differences in the SCB osteocyte expression, in both non-operated and DMM mouse joints (Fig. 4A).

Immunolabelling for sclerostin showed positive labelling in the hypertrophic chondrocytes and the SCB osteocytes in the non-operated joint of vehicle-treated mice (Fig. 4B). With DMM, there was no apparent effect on sclerostin expression by SCB osteocytes, however, there was a focal loss of sclerostin expression from the hypertrophic chondrocytes (Fig. 4B). This loss was more apparent with Bortezomib treatment in both non-operated and DMM mouse joints (Fig. 4B). These data suggest that the administration of the proteasome inhibitor Bortezomib is unable to protect against surgically induced osteoarthritis pathology, and has no effect on E11 expression in the SCB, but does diminish sclerostin expression in hypertrophic chondrocytes of the articular cartilage.

### *Applied loading does not induce articular cartilage lesions or osteophyte formation in E11 cKO mice*

As the administration of Bortezomib was unable to modify SCB E11 expression and had no effect on osteoarthritis induced by DMM, we next examined whether the conditional deletion of E11 from bone modifies susceptibility to early adaptive processes and joint vulnerability to osteoarthritis induction by a mechanically-induced post-traumatic osteoarthritis model. We have shown previously that 2 weeks of applied knee joint loading is sufficient to induce localized articular cartilage lesions in the lateral femur [14] and as such, we loaded 12-week-old cKO and WT mice and examined the lateral femur articular cartilage for load-induced lesions.

We found that, in concordance with our previous work, WT mice exhibited significant articular cartilage lesions in the lateral femur, in comparison to non-loaded limbs (p<0.01; Fig. 5A, C, E). However, 12-week-old cKO mice, in contrast, did not develop significant lateral femur lesions following 2 weeks of applied loading, in comparison to non-loaded limbs (Fig. 5B, D, E). However, no significant differences between loaded cKO and WT limbs were observed. Furthermore, osteophytes were observed by histology as expected in the loaded knee joints of all WT mice but not in loaded knee joints of E11 cKO mice (Fig. 6A & B). To identify baseline differences in articular cartilage structure which may underpin protection against load-induced lesions, we next measured articular cartilage thickness. Surprisingly, we found that the articular cartilage in the lateral femur of our E11 cKO mice (non-loaded) was significantly thinner than the WT mice (P<0.001, Fig. 6B). This was however counterbalanced by a significant increase in the thickness of the medial femur articular cartilage in the cKO mice (P<0.05, Fig. 6B). Examination of the lateral femur SCB plate surprisingly revealed no significant differences in the bone plate thickness (Fig. 6C). Similarly, no significant differences were observed in the femoral epiphyseal trabecular parameters (Fig. 6D - H). Consistent with this and with our previously published data, no significant differences were observed in osteoclast number per bone surface (Fig. 6I) [13]. Together, these data suggest that the conditional deletion of E11 from bone protects against articular cartilage lesion and osteophyte induction in response to transient joint loading.

**Discussion**

Here we reveal that the bone-specific conditional deletion of E11 in mice is protective against load induced osteoarthritis pathology. This is evidenced by the restriction of both the load-induced development of articular cartilage lesions and osteophyte formation in our E11 cKO mice. We also reveal that Bortezomib fails to exert any protection against osteoarthritis development in a surgical model of osteoarthritis (DMM). This conclusion was unexpected and opposite to our original hypothesis in which we speculated that disruption to the integrity of the osteocyte network would lead to greater osteoarthritis vulnerability.

Osteocytes are the most numerous bone cell type (>95% of bone cells) and are essential to bone structure and function. They have a unique morphology with long dendritic processes creating multicellular networks permeating the entire bone matrix [3, 21]. Historically considered passive ‘place-holders’, osteocytes have now emerged as versatile orchestrators of bone remodelling as they regulate both osteoblast (bone-forming cells) and osteoclast (bone-resorbing cells) function [22, 23]. In osteoarthritis, osteocytes are known to have a dysfunctional morphology with shorter and fewer dendritic processes [4]. As E11 is essential for the formation of osteocyte dendrites, we therefore hypothesised that the ablation of E11 from bone cells would lead to SCB thickening and exacerbated osteoarthritis pathology via a decreased osteocyte production of the bone formation inhibitor, sclerostin. However, in contrast to our expectations, we observed the opposite outcome. Our loading model used herein is non-invasive and does not induce anterior cruciate ligament rupture, thus avoiding complications that surgical methods have surrounding risk of disturbances to peri-articular tissues and disease progression. Further, whilst DMM relies on permanent, intransient destabilisation where indirect induction and progression of articular cartilage lesions are inseparable, joint loading in our model is controlled and transient, allowing direct induction of lesions and separation from progression. The data from our loading regime (6 loads in 2 weeks) thus allows examination of load-induced early osteoarthritis induction [14]. Our data indicate that less efficient osteocyte differentiation and dendrite formation, due to the hypomorphic deletion of E11, protects against the induction of osteoarthritic articular cartilage in response to transient loading episodes. Further, our data presented here suggests that a disrupted osteocyte morphology occurs in response to osteoarthritis pathology, rather than being causative. This does not however negate the possibility of E11 deficiency increasing vulnerability to other stratifications of osteoarthritis. It is also important to consider the sample size used for our WT mice as a potential limitation of this study. However, using the *in vivo* loading model, similar small group sizes have been used to resolve statistically significant load-related differences in articular cartilage lesions, surface strains by digital image correlation and osteocyte protein expression by immunolabelling [14, 24, 25]. These results indicate the very high level of reproducibility and experimental robustness of this loading model and provides us with confidence in our interpretation of our data.

Herein we also show that the bone specific deletion of E11 results in reduced articular cartilage thickness in the lateral femur. This is somewhat surprising as it is thought that a thinner articular cartilage is more susceptible to tensile strains, and therefore load induced trauma [26, 27]. Further, as both sclerostin and E11 are expressed in chondrocytes, and as no effects were seen on osteoclast activity or subchondral bone architecture, the protection to load-induced osteoarthritis afforded by E11 deletion in bone may, at least partially, reflect E11’s currently undefined role in the articular cartilage.

We have recently shown using *in vitro* osteocyte differentiation that late osteoblast E11 protein levels are regulated post-translationally by proteasome degradation and that their preservation, through use of proteasome inhibitors, such as Bortezomib, leads to the induction of an osteocyte-like morphology [9]. Bortezomib is used *in vivo* for the treatment of multiple myeloma and it is undergoing clinical trials for epithelial cancer treatment [28]. Moreover, it has been shown that Bortezomib prevents the degradation of collagen type II and the induction of MMP13 *in vitro*, thereby suggesting that it may have therapeutic effects in the context of osteoarthritis [29]. We therefore speculated that administration of the proteasome inhibitor, Bortezomib, *in vivo* would exert a protection against osteoarthritis development in an alternative surgically-induced model. We found that the administration of 1mg/kg Bortezomib, via intraperitoneal injection, to mice undergoing DMM surgery had no effect however on osteoarthritis pathology. This concentration and route of delivery has been shown previously to successively reduce proteasome 20S and mitigate histopathological manifestation of pancreatic injury in mice [18]. This is in contrast to a recent publication which showed that the administration of another proteasome inhibitor, MG132, protects against DMM-induced osteoarthritis [30]. There are many possible explanations as to why we observed these contrasting results, the most likely of which is that MG132 and Bortezomib are different types of proteasome inhibitors – MG132 is a peptide aldehyde which also inhibits certain cysteine proteinases, whereas Bortezomib is a peptide boronate inhibitor [31]. Whilst Bortezomib is currently being developed in the clinic, it would be interesting to examine the effects of other proteasome inhibitors within these subcategories to explore whether they exert modification in osteoarthritis development. It is also pertinent to consider our immunohistochemistry results, which showed that the *in vivo* administration of Bortezomib was not associated with any modification in E11 expression levels in the SCB osteocyte. The failure of *in vivo* Bortezomib administration to recapitulate its *in vitro* effects on E11 expression may indeed offer an explanation for lack of effect on osteoarthritis severity. It is nonetheless intriguing that *in vivo* Bortezomib treatment instead provoked decreased levels of E11 and sclerostin expression in articular cartilage chondrocytes – thus indicating that our Bortezomib dosing procedure was biologically effective in cartilage. The reasons for these observations require further study. It must also be borne in mind that the proteasome has manifold effects on cellular metabolic and signalling pathways, and its effects will not be limited to those we have analysed here.

The data generated here contribute to our understanding of osteoarthritis development and to our pursuit of a disease-modifying treatment. We have shown that the clinically-relevant drug Bortezomib, was not found in this study to have any therapeutic potential in a surgical model of osteoarthritis. We have however shown that the precise control of E11 is crucial in SCB function in osteoarthritis and that the regulatory networks controlling E11 osteocyte expression are more complex *in vivo* than they are *in vitro.* Furthermore, the data presented here offer further support for the role of cartilage: bone interactions in the development of osteoarthritis.

**Conclusions**

Overall, these data suggest that an intact osteocyte network in the SCB contributes to the development of mechanically-driven osteoarthritis. Further, the data presented here indicate that the molecular pathways that preserve the osteocyte network, such as those driven by E11, may be targeted to limit osteoarthritis pathogenesis.

**List of abbreviations**

BV/TV – bone volume/tissue volume; cKO – conditional knockout; DMM - destabilisation of the medial meniscus; LF – lateral femur; LT – lateral tibia; MF – medial femur; MT – medial tibia; microCT – micro-computed tomography; ROI – region of interest; SCB – subchondral bone; SCB Th. - subchondral bone thickness; Tb. BV/TV - trabecular bone volume/tissue volume; Tb. N. - trabecular number; Tb. Pf. - trabecular pattern factor; Tb. Sp. - trabecular separation; Tb. Th. – trabecular thickness; WT – wild-type;

**Declarations**

*Ethics approval and consent to participate*

All experimental protocols were approved by Roslin Institute's Animal Users Committee and the animals were maintained in accordance with UK Home Office guidelines for the care and use of laboratory animals. The human samples were obtained with written patient consent and all procedures with ethical approval by NHS Lothian. The collection, storage, and subsequent use of human tissues are regulated in Scotland by The Human Tissue Act (Scotland) 2006. Canine osteoarthritis samples were collected with informed written consent from the owner and ethical approval for their collection and use given by the Veterinary Ethical Review Committee of the University of Edinburgh (VERC; approval 23/12). We obtained written informed consent to use the osteocalcin-cre mice in our study from Dr Thomas Clemens.

*Consent to publish*

Not applicable

*Availability of data and material*

The datasets used and/or analysed during the current study are available from the corresponding author on reasonable request.

*Competing interests*

The authors declare that they have no competing interests.

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*Authors' contributions*

Conception and design of the study: all authors, Acquisition of data: KAS, AT, SD, EI, BJ, AKA, Interpretation of data: KAS, EI, DNC, DJB, AAP, CF, Drafting the manuscript: KAS, AAP, CF, Revising the manuscript and final approval, and agreement to be accountable for all aspects of the work: all authors.

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**Figure legends**

**Figure 1.** Immunohistochemical labelling for E11 in (**A**) DMM and non-operated C57/BL6 mice (**B**) human osteoarthritis samples from patients undergoing total knee replacement (**C**) canine osteoarthritis samples from the medial coronoid process of the elbow. Images are representative of n>3. Arrows are representative of E11 positive osteocytes. Scale bar = 300µm.

**Figure 2.** **(A)** Weights of mice treated with Bortezomib or vehicle for days post DMM surgery **(B)** Maximum OARSI score in the medial and lateral compartments of the left (contralateral control) knee joint of Bortezomib and vehicle treated mice (**C)** Maximum OARSI score in the medial and lateral compartments of the right (DMM) knee joint of Bortezomib and vehicle treated mice (**D)** Mean OARSI score in the medial and lateral compartments of the left non-operated (contralateral control) knee joint of Bortezomib and vehicle treated mice (**E)** Mean OARSI score in the medial and lateral compartments of the right (DMM) knee joint of Bortezomib and vehicle treated mice (**F)** Representative histology images of articular cartilage lesions (arrows) in the medial joint compartment. Data are presented as mean ± S.E.M (n=8/group).

**Figure 3.** MicroCT analysis of the epiphyseal region of the medial tibiain DMM-operated and non-operated controls **(A)** subchondral bone thickness (SCB Th.) **(B)** trabecular bone volume/tissue volume (Tb. BV/TV) **(C)** trabecular number (Tb. N.) **(D)** trabecular thickness (Tb. Th.) **(E)** trabecular separation (Tb. Sp.) **(F)** trabecular pattern factor (Tb. Pf.). MicroCT analysis of the epiphyseal region of the lateral tibia in DMM-operated and non-operated controls **(G) s**ubchondral bone thickness (SCB Th.) **(H)** trabecular bone volume/tissue volume (Tb. BV/TV) **(I)** trabecular number (Tb. N.) **(J)** trabecular thickness (Tb. Th.) **(K)** trabecular separation (Tb. Sp.) **(L)** trabecular pattern factor (Tb. Pf.). Data are presented as mean ± S.E.M (n=8/group). P<0.05\*

**Figure 4. (A)** E11 immunohistochemistry in the left non-operated (contralateral control) and right (DMM) knee joints of vehicle and Bortezomib treated mice **(B)** sclerostin immunohistochemistry in the left non-operated (contralateral control) and right (DMM) knee joints of vehicle and Bortezomib treated mice. Arrows are representative of E11 positive osteocytes and asterisks indicate positive E11 immunolabelling in the chondrocytes.

**Figure 5. (A)** Mean OARSI score in the lateral femur of control and loaded knee joints of WT mice. **(B)** Mean OARSI score in the lateral femur of control and loaded knee joints of cKO mice. **(C)** Maximum OARSI score in the lateral femur of control and loaded knee joints of WT mice. **(D)** Maximum OARSI score in the lateral femur of control and loaded knee joints of cKO mice. **(E)** Representative histology images of articular cartilage lesions (arrows) in the lateral femur. Data are presented as mean ± S.E.M (n>3/group). P<0.05\*; P<0.01\*\*.

**Figure 6.** **(A)** Histology images of osteophyte formation (arrows) in loaded joints from WT mice **(B)** Articular cartilage thickness (LF – lateral femur, LT – lateral tibia, MF – medial femur, MT – medial tibia). MicroCT analysis of the epiphyseal region of the lateral femur **(C)** subchondral bone plate thickness (SCB Th.) **(D)** epiphyseal trabecular bone volume/tissue volume (Tb. BV/TV) **(E)** epiphyseal trabecular number (Tb. N.) **(F)** epiphyseal trabecular thickness (Tb. Th.) **(G)** epiphyseal trabecular separation (Tb. Sp.) **(H)** epiphyseal trabecular pattern factor (Tb. Pf.). (**I**) Number of osteoclasts / bone surface. Data are presented as mean ± S.E.M (n>3/group). P<0.05\*; P<0.001\*\*\*