Characterisation of growth plate dynamics in murine models of osteoarthritis

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Abstract

Background: The purpose of this study was to investigate growth plate dynamics in surgical and loading murine models of osteoarthritis, to understand whether abnormalities in these dynamics predict those at risk of osteoarthritis.

Methods: 8-week-old C57BL/6 male mice underwent destabilisation of medial meniscus (DMM) ($n=8$) surgery in right knee joints. Contralateral left knee joints had no intervention (controls). In 16-week-old C57BL/6 male mice ($n=6$), osteoarthritis was induced using non-invasive mechanical loading of right knee joints with peak force of 11N. Non-loaded left knee joints were internal controls. Chondrocyte transiency in tibial articular cartilage and growth plate was examined by histology and immunohistochemistry. Tibial subchondral bone parameters were measured using microCT and correlated to GP bridging.

Results: Higher expression of chondrocyte hypertrophy markers; Col10a1 and MMP13 were observed in tibial articular cartilage chondrocytes of DMM and loaded mice. In tibial growth plate, Col10a1 and MMP13 expressions were widely dispersed in a significantly enlarged zone of proliferative and hypertrophic chondrocytes. 3-dimensional quantification revealed enriched growth plate bridging and higher bridge densities in medial compared to lateral tibiae of DMM and loaded knee joints of the mice. Growth plate dynamics were associated with increased subchondral bone volume fraction (BV/TV; %) in medial tibiae of DMM and loaded knee joints and epiphyseal trabecular bone volume fraction in medial tibiae of loaded knee joints.

Conclusions: The results confirm associations between aberrant chondrocyte hypertrophy marker expression and osteoarthritis pathology in a surgical and loaded murine model of osteoarthritis. Spatial variations in growth plate bridging formation revealed accelerated

45 cartilage-bone transitions which may contribute to anatomical variation in vulnerability to

46 osteoarthritis development in these models.

47 **Keywords:** osteoarthritis, cartilage, growth, chondrocyte, murine model

Background

Osteoarthritis (OA) is a chronic musculoskeletal disease and a leading cause of disability and

major healthcare costs in the world. It is estimated that worldwide 10% of men and 18% of

women aged over 60 years have symptomatic OA (1) and, therefore, the predicted increase in

the ageing population and longevity will result in a greater occurrence of the disease. OA is a

complex disease in which the pathogenesis, cellular and molecular mechanisms of initiation

and progression are not completely understood. It is characterised by progressive loss of

articular cartilage (AC), formation of osteophytes, subchondral bone (SCB) sclerosis,

synovial proliferation, inflammation and lax tendons. These can ultimately lead to a loss of

joint function, pain, reduced mobility and disability (2). Despite the significant healthcare and

economic burden there are few non-invasive therapies available to patients. Therefore,

understanding the pathogenesis of OA and defining the molecular mechanisms underpinning

AC degeneration can lead to the development of successful targeted and effective disease-

modifying treatments.

Primary OA is described as naturally occurring OA affecting one joint (localised) or three or

more joints (generalised), while secondary OA is associated with various causes and risk

factors leading to the disease including trauma, obesity, diabetes, metabolic bone and

congenital disorders (3). AC degeneration is one of the main hallmarks of OA and previous

research has largely sought to identify mechanisms underpinning its deterioration. Fully

developed, uncalcified AC is populated by a single resident cell chondrocytes, which

maintain a stable phenotype characterised by small cell size and expression of tenascin-C (4).

The inherent stability of AC chondrocytes ensures that dynamic events are restricted to assure

lifelong articular integrity and healthy joint function. In contrast, epiphyseal growth plate

(GP) chondrocytes have a transient phenotype to ensure long bone development

(endochondral ossification) and growth. GP chondrocytes undergo a differentiation sequence

of proliferation, maturation and hypertrophy. The final stage of chondrocyte hypertrophy

enables mineralisation of the cartilage extracellular matrix, vascular invasion and subsequent

replacement of the mineralised cartilage anlagen with bone (5). These processes are coupled,

however, with sexual maturation the human GP undergoes progressive narrowing as bony

bridges form and span its width. This ultimately leads to complete GP closure and cessation

of human growth. Indeed, in humans the longitudinal bone growth stops with the onset of

puberty, the metaphysis then fuses with the epiphysis and the GP disappears (6). In mice,

longitudinal bone growth does not cease at sexual maturity instead it slows dramatically at

puberty, but the GPs do not completely fuse and disappear (7).

We have previously shown that in the STR/Ort mouse, a naturally occurring OA murine

model, AC chondrocytes transform from their inherently stable phenotype to a transient one,

characteristic of the chondrocytes in the GP. This was confirmed by immunolabelling for

chondrocyte hypertrophy markers; type X collagen (Col10a1) and matrix metalloproteinase

13 (MMP13)(8). Further, we revealed accelerated longitudinal bone growth, aberrant

expression of GP markers (Col10a1 and MMP13) and increased GP chondrocyte maturation

in these mice. Consistent with this, using a novel synchrotron computed tomography method

we revealed enriched GP bone bridging in STR/Ort mouse tibiae indicative of advanced GP

closure which may underpin OA (8,9).

Despite this, interlinks between the differing chondrocyte phenotypes in the AC and the GP,

and the contribution that the GP may play in underpinning OA vulnerability is not yet

understood. The aim of this study was to investigate whether surgically (destabilisation the

medial meniscus [DMM]) or non-surgically (mechanical loading) induced OA onset in

C57BL/6 mice is linked to altered GP dynamics. Indeed, understanding this will inform

strategies for maintaining musculoskeletal health in ageing by potentially identifying whether

GP dynamics may predict who is at risk of OA in later life and ultimately developing targeted

OA treatments.

Methods

Animals

Male C57BL/6 wild type mice at 7 weeks of age (young adult) were obtained from Charles

River Laboratories Inc. (Margate, UK). The mice were acclimatised to their surroundings for

seven days. All mice were allowed free access to water and maintenance diet ad libitum

(Special Diet Services, Witham, UK) in a 12-hour light/dark cycle at a room temperature of

21 ± 2^0C and relative humidity of $55 \pm 10\%$.

The destabilisation of the medial meniscus (DMM)

A group of eight 8-week old wild type C57/BL6 male mice underwent DMM surgeries to

induce OA-like changes in the right knee joints under isoflurane-induced anaesthesia. We

chose not to performed SHAM (placebo) surgery on the left contralateral knee of the animals

based upon animal welfare grounds since it was previously shown that there is no difference

in OA scores between SHAM-operated and non-operated knee joints (10,11). Animals were

monitored for unexpected adverse effects to reduce suffering and distress. Following

transection of the medial meniscotibial ligament (MMTL) to destabilise the medial meniscus,

the skin was closed and anaesthesia reversed (10). Eight weeks later, mice were culled by

exsanguination and confirmation of death by cervical dislocation. The knee joints of all the

mice were dissected, fixed in 4% paraformaldehyde for 24 hours at 4°C, and then stored in

70% ethanol.

In vivo loading of the knee joint

The right knee joints of 16-week-old wild type C57BL/6 male mice ($n = 6$) were subjected to

non-invasive, dynamic axial mechanical loading under the isofluorane-induced anaesthesia

(liquid isofluorane was vaporised to a concentration of 4% and maintained at a concentration

of 2% with oxygen) for 7 min/day, 3 alternate days a week for 2 weeks according to the

protocols described in the previous studies (12,13). The left knee joints were non-loaded

internal controls in these animals. Briefly, using a servo-electric materials testing machine

(Electroforce 3100, Bose, UK), axial compressive loads were applied through the right knee

joint via customised concave cups which held the knee and ankle joints flexed and the tibiae

vertically.

The tibia was held in place by continuous static preload of 0.5N onto which dynamic loads

were superimposed in a series of 40 trapezoidal shaped waveform cycles with steep up and

down ramps and a peak force of 11N for 0.05 seconds (0.025 seconds rise and fall time; 9.9

seconds baseline hold time between periods of peak loading). The right and left knees were

dissected 3 days after the final loading episode. Mice were culled by exsanguination and

confirmation of death by cervical dislocation. Knee joints were fixed in 4%

paraformaldehyde for 24 hours at 4°C before being stored in 70% ethanol.

Micro-computed (microCT) tomography and 3-dimensional (3D) bridging analysis

Scans were performed with an 1172 X-Ray microtomograph (Bruker MicroCT, Kontich,

Belgium) to evaluate the SCB and GP bridging. High-resolution scans with an isotropic voxel

size of 5 µm were acquired (50 kV, 200 µA, 0.5 mm aluminium filter, 0.6° rotation angle).

The projection images were reconstructed and binarised with a threshold of 0 to 0.16, ring

artefact reduction was set at 10 and beam hardening correction at 0% using the SkyScan

NRecon software package (v1.6.9.4, Bruker MicroCT). The images then were realigned

vertically using DataViewer software (v1.5.1.2 64-bit, Bruker MicroCT) to ensure similar

orientation for analysis. Hand-drawn regions of interests (ROI) of the SCB and epiphyseal

trabecular bone for each tibial lateral and medial compartments were selected (14). The

structural parameters of tibial SCB plate and epiphyseal trabecular bone were calculated

using 3D algorithms of SkyScan CTAn software (Bruker MicroCT) including SCB (SCB

BV/TV; %) and trabecular bone volume fraction (Tb. BV/TV; %) and correlated to GP

bridging analysis using a 3D synchrotron-computed tomography quantification method as

previously described (15). Briefly, microCT scans of the tibiae were segmented using a

region-growing algorithm within the Avizo® (V8.0, VSG, Burlington, VT, USA) software.

The central points of each bony bridges were identified and projected on the tibial joint

surface. The distribution of the areal number density of bridges (N, the number of bridges per

256 µm × 256 µm window; $d = m/V$) is then calculated and superimposed on the tibial joint

surface (each bridge has a colour that represents the areal number density at the bridge

location). The SCB plate and epiphyseal trabecular bone thickness (Th; mm) was determined

and colour-coded thickness images were generated using the Avizo® software(15).

Histological analysis

The left and right knee joints of all the mice were decalcified in 10%

ethylenediaminetetraacetic acid (EDTA) solution, wax-embedded at Leica EG1160 Tissue

Embedding Station and 6 µm coronal sections cut using Leica RM2135 manual microtome.

Immunohistochemistry

Immunohistochemical analysis of chondrocyte transiency markers in tibial AC and GP was

performed on 6 µm coronal sections using anti-matrix metalloproteinase 13 (anti–MMP13)

(1:200 dilution; Abcam) or anti-collagen type X (anti-Col10a1) (1:100 dilution; Abcam)

antibodies. As a control, an equal concentration of rabbit IgG was used. For

immunohistochemical localisation of MMP13 and Col10a1, 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 0.3% H_2O_2 in methanol

(Sigma) for 30 min at room temperature. The Vectastain ABC universal detection kit (Vector

Laboratories, Peterborough, UK) was used to detect the biotinylated secondary antibody

(Anti-Mouse IgG Reagent) after incubation for 30 min at room temperature according to the

manufacturer's instructions. Diaminobenzidine (DAB) solution used to detect the location of

antigens. The sections were finally dehydrated, counterstained with haematoxylin and

mounted in DePeX. All sections to be compared were immunolabelled at the same time to

standardise conditions and minimise any differences in antibody incubation times.

Growth plate (GP) zone analysis

The stained sections from the joints of 4 individual mice from each experimental group were

used to measure the width of the GP proliferating and hypertrophic zones, as well as the total

GP width, measured at 10 different points along the length of the GP in tibiae, using a light

microscope and ImageJ software.

Statistical analysis

All analyses were performed with GraphPad Prism software 6.0f version (GraphPad Inc, La

Jolla, CA, USA) using a two-sided 0.05 level of significance. All analyses were conducted

blindly to minimise the effects of subjective bias. The results were presented as the mean ±

standard error of the mean (SEM). The Normal distribution of data was assessed using the

Shapiro-Wilk normality test. For comparing two groups (experimental with control, or medial

with the lateral compartment of tibiae), two-tail Student's *t*-test (paired or unpaired) was

used. For comparing more than two groups, two-way ANOVA (analysis of variance) was

used with Tukey post-hoc test.

Results

Transient chondrocyte behaviour in the tibial AC of DMM and loaded C57BL/6 young adult

male mice

We first sought to confirm whether loss of AC in C57BL/6 mice with surgically and loading

induced OA was associated with the expression of markers of transient chondrocyte

phenotype. Immunohistochemistry analysis showed higher expression levels of well-

established chondrocyte hypertrophy markers; Col10a1 and MMP13, observed in tibial AC of

C57BL/6 mice that have undergone DMM surgery or mechanical loading compared with

non-operated and non-loaded control left tibiae, respectively (Fig. 1a, and b). The expression

pattern of Col10a1 was largely restricted to hypertrophic chondrocytes in the uncalcified zone

of the AC of unaffected condyles of non-operated (Fig. 1a) and non-loaded (Fig. 1b) mouse

left joints as expected (16). Whereas, the immunolabeling of Col10a1 was more widespread

throughout the extracellular matrix (ECM) of the AC in affected right joints of DMM (Fig.

1a) and loaded (Fig. 1b) mice. Similarly, immunohistochemistry analysis showed positive

MMP13 labelling in both superficial and deep articular chondrocytes in the right joints of

DMM and loaded C57BL/6 male mice compared to the control knee joints (Fig. 1a and b).

These findings confirm an aberrant deployment of transient chondrocytes in uncalcified AC.

Dysfunctional GP morphology in DMM and loaded tibiae of C57BL/6 young adult male mice

In the tibial GP of mice with surgically and loading induced OA, Col10a1 expression was

more greatly and widely dispersed throughout the zones of proliferative and hypertrophic

chondrocytes compared with their controls (Fig. 2a and b). Indeed, immunolabeling for

Col10a1 revealed the expected localisation in the GP of non-operated and non-loaded mouse

tibiae, limited primarily to the hypertrophic zone and underlying adjacent metaphyseal bone

(Fig. 2a and b). This disrupted the distribution of a GP zone marker was also evident for

MMP13 in the GP of DMM and loaded C57BL/6 mouse tibiae compared to their controls

(Fig. 2a and b).

GP zone analysis revealed significantly enlarged proliferative and hypertrophic zones of

chondrocytes in both DMM (p=0.002 and p<0.0001, respectively) and loaded (both

p<0.0001) tibiae of C57BL/6 mice compared to their controls, and significantly increased

total GP width (both p<0.0001) (Fig. 3). Together, the results may indicate associations

between dysfunctional GP morphology and marker expression, and OA development.

Associations between GP bridging in DMM and loaded tibiae of C57BL/6 young adult male

mice and OA development

To further correlate aberrant longitudinal GP dynamics, GP bridging and OA development in

these C57BL/6 young adult male mice, we used our newly developed 3D method to quantify

bony bridges across the tibial epiphysis of DMM and loaded mice (Fig. 4). 3D quantification

revealed a significantly higher number of GP bridges in medial compared to lateral tibiae that

underwent DMM surgeries (196 $\pm$ 33 versus 306 $\pm$ 32; p=0.03) (Fig. 4b and e). This

significant difference was not observed in non-operated tibiae (254 $\pm$ 32 versus 326 $\pm$ 32;

p=0.14) (Fig. 4a and e). Similarly, significantly enriched GP bridging was evident in the

medial compartment of loaded tibiae in comparison to the lateral compartment (539 $\pm$ 36

versus 745 $\pm$ 18; p=0.0009) (Fig. 4d and e), and in those of non-loaded tibiae, although less

pronounced than in the loaded right knee joints (596 $\pm$ 22 versus 726 $\pm$ 44; p=0.04) (Fig. 4c

and e). However, no significant differences in GP bridge numbers and densities were

observed between interventions (DMM versus non-operated, and loaded versus non-loaded)

at this time point in either the medial or lateral compartment. Anatomical variations were

observed however with clusters forming in the medial anterior of loaded tibiae versus those

more in the central and posterior of non-loaded tibiae (Fig. 4c and d). Similarly, in the DMM

tibiae, GP bridges were more widespread across the tibiae than in non-operated which were

predominantly observed around the periphery (Fig. 4a and b). These results were consistent

with the areal bridge density analysis. The mean areal bridge densities were significantly

greater in medial compared to the lateral compartment of DMM (8.2 ± 0.9 versus 10.3 ± 0.9;

p=0.01) and loaded tibiae (16.3 ± 0.6 versus 20.6 ± 0.5; p<0.0001) (Fig. 4f). However, no

significant differences in the mean areal bridge densities were observed between

interventions (DMM versus non-operated, and loaded versus non-loaded).

MicroCT analysis and 3D visualisation of SCB plate and epiphyseal trabecular bone

To establish whether aberrant GP dynamics are associated with the SCB plate and epiphyseal

trabecular bone abnormalities after the DMM surgery and mechanical loading in our mice,

we performed microCT analysis and determined local thickness of SCB plate and epiphyseal

trabecular bone. The SCB plate volume fraction (SCB BV/TV) was significantly higher in

medial compared to lateral compartment of DMM and non-operated tibiae (DMM: SCB

BV/TV 29.4 ± 3.1% versus 36.7 ± 4.5%, p=0.03, non-operated: SCB BV/TV 35.7 ± 0.7%

versus 42.6 ± 1.5%, p=0.004) (Fig. 5a). No significant differences were observed between

DMM and non-operated tibia (Fig. 5a), or in the epiphyseal trabecular bone volume fraction

(Tb. BV/TV; Fig. 5b). Conversely, in the loaded and non-loaded tibiae, epiphyseal trabecular

bone volume fraction (Tb. BV/TV) was significantly increased in the medial compared to

lateral compartment (loaded: Tb. BV/TV 60.7 ± 1.8% versus 73.4 ± 2.8%, p=0.005, non-

loaded: Tb. BV/TV 61.4 ± 2.2% versus 77.4 ± 2.5%, p=0.0005). No significant differences

were observed between interventions (Fig. 5d). The SCB plate volume fraction (SCB

BV/TV) was significantly higher in medial compared to lateral compartment of loaded tibiae

(40.1 ± 1.7% versus 47.1 ± 1.6%, p=0.02), but not that of non-loaded tibiae (42.1 ± 1.1%

versus 45.8 ± 1.6%, p=0.34) or between interventions (Fig. 5c). Colour-coded SCB plate

thickness analysis revealed anatomical variation in SCB plate thickness between DMM and

non-operated, and loaded and non-loaded plates (Fig. 6a and 6b). Differences in the non-

loaded and loaded SCB plates were particularly apparent (Fig. 6b), and correlated with the

clusters of higher density of GP bridges previously observed (Fig. 4c and d). Epiphyseal

trabecular bone thickness was not significantly altered in response to either invasive or non-

invasive intervention (data not shown; Fig. 6c and d).

Discussion

This study reveals altered GP dynamics in both surgical (DMM) and non-invasive loading

murine *in vivo* models of OA, which with findings described above may indicate an

endochondral defect in AC and GP cartilage in these mouse models of OA. Our data show

changes in AC of the knee joints of these mice consistent with the aberrant deployment of

hypertrophic chondrocytes. This is associated with disrupted GP morphology, increased GP

chondrocyte differentiation indicated by widespread expression of chondrocyte hypertrophy

markers and increased GP zone widths. Moreover, we have discovered enriched GP bony

bridging indicative of premature GP fusion and accelerated growth cessation in the medial

compartment of tibiae of these mice. These bridging events are spatially correlated with

increase in SCB thickness. These data reveal that altered GP dynamics and spatial differences

in GP bridging may contribute to an anatomical variation in vulnerability to OA development

in surgical and loaded murine models of OA.

The STR/ort OA murine model is predisposed to developing spontaneous idiopathic OA

whilst its nearest available parental strain, the CBA mouse, has a very low susceptibility

which makes them effective controls for the studies (17). We have previously shown that

aberrant deployment of transient chondrocyte behaviour, consistent with re-initiation of

endochondral processes, occurs in uncalcified AC of STR/ort mouse knee joints compared to

CBA controls (8). Here we extend these studies to look at the expression of transient markers

in other stratifications of OA. Surgically induced DMM model is widely used for target

validation studies or evaluation of the pathophysiological roles of many molecules in OA.

Following DMM, medial displacement of the medial meniscus in a mouse knee joint provides

a smaller area to transmit the weight-bearing forces and leads to an increased local

mechanical stress (18). Whereas, in a cyclic AC tibial compression model, the non-invasive

dynamic mechanical loading applied to the mouse tibia through the knee and ankle joints,

modifies AC structure locally through a mechanoadaptive homeostatic response contributing

to OA development (19). Our findings in both surgical and non-invasive loading C57BL/6

mouse models are consistent with the evidence of the role of chondrocyte phenotype

alterations in OA pathology. Hypertrophic chondrocytes in the calcified cartilage and GP of

the healthy joints express Col10a1 (17,19). The calcified cartilage acts to protect the

uncalcified AC through maintaining its ECM in an unmineralised state and the stability of the

AC. However, hypertrophic differentiation of these chondrocytes contributes to AC matrix

degradation, calcification and vascular invasion resulting in the demise of the AC (23).

Consistent with this, the expression of Col10a1, as examined using immunohistochemistry,

has been observed throughout AC in the joints of our both DMM and loaded mice. Further,

the higher expression level of another marker of chondrocyte hypertrophy; MMP13 has been

detected in superficial and uncalcified chondrocytes in the AC of DMM and loaded mice

compared to their non-operated and non-loaded left knee joints. Indeed, cartilage degradation

observed in OA has been attributed to an elevated production of proteolytic enzymes among

which MMP13 has a major role (24,25). Studies using transgenic mice deficient in catabolic

transcription factors that induce hypertrophic differentiation revealed that animals were

protected against surgically and chemically induced OA further highlighting the role of

transient chondrocytes in AC degradation (26,27).

The results of the present study indicate aberrant widespread expression of Col10a1 and

MMP13 in the GP of DMM and loaded tibiae of C56BL/6 mice, compared to the GP of non-

operated and non-loaded control knees. Indeed, the histomorphometric analysis also revealed

that these mice display a significantly enlarged zone of proliferative and hypertrophic

chondrocytes, compared with the chondrocytes in the GP of control tibiae of both models.

Similarly, a significantly increased cumulative GP width was observed in these models.

Longitudinal bone growth is determined by the modifying number of chondrocytes in the

proliferative zone of GP, rate of their proliferation, the extent of chondrocyte hypertrophy

and controlled synthesis and degradation of ECM throughout the GP (28). Altered growth

rate and mechanical modulation of GP function appear to result from complex interactions of

changes in the states of these chondrocytes, as does the rate of GP closure due to the

formation of bone bridges forming and spanning the width of the GP.

Our report of increased GP chondrocyte dynamics in these DMM and loaded C57BL/6 mice

is further strengthened by our data acquired using a 3D quantification method of bony

bridging across the tibial epiphysis, which showed premature GP closure in the medial

compartment of the tibiae in all mice examined, regardless of intervention. This was

indicated by significantly enriched spatial localisation of GP bone bridging clustering and

number in the medial compared to the lateral compartments of DMM mice compared to non-

operated mice. This is consistent with our previous work in the STR/Ort mouse model of

spontaneous OA in which we observed similar spatial variations. We postulate that the

formation of these bridges may be accelerated by local factors like instigating altered

mechanoadaptive response and that these spatial variations in GP bridging may disclose the

anatomical vulnerability to OA. These findings are supported by the previous studies

suggesting associations between local mechanical stress caused by medial displacement of

the medial meniscus (29) or cyclic tibial AC compression (19) and GP function in C57BL/6

mice. They are further supported by our previous work in which we revealed, by finite

element modelling, that GP bridges act to dissipate stresses upon loading to the overlying

SCB and thus suggest that this contributes to OA seen in these models. These studies have,

for the first time, shown that accelerated GP closure is indicative of modified growth

dynamics in the right tibiae of both surgical and loading murine models of OA. This is

consistent with our microCT analysis of spatial variations in SCB plate thickening in these

animals, and its correlation to GP bridges. We observed statistically significant changes in

subchondral bone parameters in medial compared to lateral compartment including

subchondral bone volume fraction in the tibiae of both DMM and loaded mice and epiphyseal

trabecular bone volume fraction in the tibiae of loaded mice. It was previously shown that

SCB thickening may be intensified by adjacent AC lesions following mechanical trauma thus

indicating a special link between changes in SCB architecture and AC lesions in OA

pathology (14).

The limitation of this study is that we only examined the GP at one specific time point for

each model and therefore more time points are required to understand how these GP bridges

temporally affect SCB changes and OA pathology. With its controllability, the intermittent

non-invasive mechanical loading model will allow in future to distinguish between short- and

long-term effects of various cyclic loading regimens on SCB and trabecular bone parameters

as well as AC integrity, GP dynamics and correlate these to initiation and progression of

human OA. Indeed, it is known that in the loading model, the short-term intervention is not

sufficient to induce significant changes in subchondral thickness and thus longer intervention

times for both the DMM and loading models may also prove useful in pursuit of

understanding these relationships (14).

Conclusions

Our studies indicate that accelerated GP chondrocyte dynamics and bridging events may

contribute to OA pathology in both surgical and loading C57BL/6 mouse OA models and that

similar osteoarthritic pathological changes happen in different *in vivo* models of secondary

(post-traumatic) OA (18) through inducing direct (DMM) or indirect (mechanical loading)

injuries to the joints. The GP bony bridging analysis may signify accelerated cartilage-bone

transition and growth cessation in these affected bones advancing our understanding of GP

closure mechanisms and how these contribute to the health of the joint. Further, our work

yields more insights into the changes in the micro-mechanical environment of the GP and

cells, specifically chondrocytes, within the GP.

Nonetheless, whether modified GP trajectories are prevalent in human OA is yet to be

defined and studies determining associations between GP dynamics during adolescence and

OA development in human patients would allow elucidation of the pathogenesis of OA and

will ultimately enable the development of novel and specific therapeutic interventions.

List of abbreviations

AC, articular cartilage; Col10a1, type X collagen; DAB, Diaminobenzidine; DMM,

destabilisation of medial meniscus; ECM, extracellular matrix; GP, growth plate; MMP13,

matrix metalloproteinase 13; MMTL, meniscotibial ligament; OA, osteoarthritis; SCB,

subchondral bone.

Declarations

Ethics approval and consent to participate

All procedures complied with the United Kingdom Animals (Scientific Procedures) Act 1986

and were approved by The University of Edinburgh Roslin Institute's Animal Users and

Research Ethics Committees.

Consent for publication

Not applicable

Availability of data and materials

The datasets used and/or analysed during the current study are available from the

corresponding authors on reasonable request.

Competing interests

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Authors' contributions

All authors contributed to the study design, analysis and interpretation of the data, drafted,

critically reviewed, edited and approved the version of the manuscript for publication.

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Figure legends

Fig. 1. Immunohistochemical labelling in the tibial articular cartilage of mouse knee

joints

Immunohistochemical labelling for type X collagen (Col10a1) and matrix metalloproteinase

(MMP13) in the articular cartilage of non-operated and DMM (**a**), or non-loaded and

mechanically loaded (**b**) knee joints of C57BL/6 male mice. Arrows indicate examples of

positive staining in the tibiae. Images are representative of results in 3 individual mice. Scale

bar = 200µm

Fig. 2. Immunohistochemical labelling in the tibial growth plate of mouse knee joints

Immunohistochemical labelling for type X collagen (Col10a1) and matrix metalloproteinase

(MMP13) in the growth plate of DMM and non-operated (**a**), or mechanically loaded and

non-loaded (**b**) knee joints of C57BL/6 male mice. Arrows indicate examples of positive

staining in the tibiae. Images are representative of results in 3 individual mice. Scale bar =

200µm. PZ = proliferative zone; HZ = hypertrophic zone

Fig. 3. Measurement of the growth plate zone width of mouse knee joints

Growth plate zone width of non-operated and DMM, or non-loaded and loaded knee joints of

C57BL/6 male mice. Ten measurements per section were obtained along the length of the

growth plate of the tibiae (n=4 mice for each experimental group). PZ = proliferative zone;

HZ = hypertrophic zone **** $p<0.0001$

Fig. 4. Bridge location and bridge areal densities across the tibial growth plate of mice

Location and areal densities of bridges across the growth plate projected on the medial (M)

and lateral (L) tibial joint surface in non-operated (**a**), DMM (**b**), non-loaded (**c**) and loaded

(**d**) tibiae of mice at 16 and 18 weeks of age, number of bridges in lateral and medial tibiae of

non-operated compared to DMM and non-loaded compared to loaded tibiae of mice (**e**),

Areal density (d) of bridges in medial compared to lateral tibiae, defined as the number of

bridges per 256 mm x 256 mm window of non-operated and DMM or non-loaded and loaded

knee joints (F). Bars represent mean ± SEM. Group sizes were $n = 8$ for non-operated and

DMM-operated mice and $n = 6$ for non-loaded and loaded mice. * $p<0.05$ ** $p<0.01$ ***

$p<0.001$ **** $p<0.0001$

Fig. 5. MicroCT analysis of epiphyseal region of medial and lateral tibiae of mice

MicroCT analysis of the epiphyseal region of the lateral and medial tibiae in non-operated

controls and DMM-operated knee joints subchondral bone volume fraction (SCB BV/TV) (**a**)

and epiphyseal trabecular bone volume fraction (Tb. BV/TV) (**b**). MicroCT analysis of the

epiphyseal region of the medial and lateral tibiae in non-loaded controls and loaded knee

joints subchondral bone volume fraction (SCB BV/TV) (**c**) and epiphyseal trabecular bone

volume fraction (Tb. BV/TV) (**d**). Bars represent mean ± SEM. Group sizes were $n = 8$ for

non-operated and DMM-operated mice and $n = 6$ for non-loaded and loaded mice. * $p<0.05$

** $p<0.01$ *** $p<0.001$

Fig. 6. Colour-coded local thickness of tibial subchondral bone plate and epiphyseal

trabecular bone of mice

Representative colour coded images of lateral and medial subchondral bone plate thickness of

non-operated and DMM-operated tibiae of mice (**a**). Representative colour coded images of

lateral and medial subchondral bone plate thickness of non-loaded and loaded tibiae of mice

(**b**). Representative colour coded images of epiphyseal trabecular bone thickness of non-

operated and DMM-operated (**c**), or non-loaded and loaded tibiae of mice (**d**).

a Non-operated | DMM

Col10a1

MMP13

PZ · HZ

b Non-loaded | Loaded

Col10a1

MMP13

PZ · HZ

a

Non-operated

Lateral   Medial

DMM

Lateral   Medial

c

Non-operated

Lateral   Medial

DMM

Medial   Lateral

b

Non-loaded

Lateral   Medial

Loaded

Lateral   Medial

d

Non-loaded

Lateral   Medial

Loaded

Medial   Lateral