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The effect of Tlr4 and/or C3 Deficiency and of Neonatal Gene Therapy on Skeletal Disease in Mucopolysaccharidosis VII mice

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Abstract

Mucopolysaccharidosis (MPS) VII is a lysosomal storage disorder caused by the deficiency of the enzyme β -glucuronidase (*Gusb*^{-/-}) and results in glycosaminoglycan (GAG) accumulation. Skeletal abnormalities include stunted long bones and bone degeneration. GAGs have been hypothesized to activate toll-like receptor 4 (Tlr4) signaling and the complement pathway, resulting in upregulation of inflammatory cytokines that suppress growth and cause degeneration of bone. *Gusb*^{-/-} mice were bred with *Tlr4*- and complement component 3 (*C3*)-deficient mice, and the skeletal manifestations of the doubly- and triply-deficient mice were compared to those of purebred *Gusb*^{-/-} mice. Radiographs showed that purebred *Gusb*^{-/-} mice had shorter tibias and femurs, and wider femurs, compared to normal mice. No improvement was seen in *Tlr4*, *C3*, or *Tlr4/C3*-deficient *Gusb*^{-/-} mice. The glenoid cavity and humerus were scored on a scale from 0 (normal) to +3 (severely abnormal) for dysplasia and bone irregularities, and the joint space was measured. No improvement was seen in *Tlr4*, *C3*, or *Tlr4/C3*-deficient *Gusb*^{-/-} mice, and their joint space remained abnormally wide. *Gusb*^{-/-} mice treated neonatally with an intravenous retroviral vector (RV) had thinner femurs, longer legs, and a narrowed joint space compared with untreated purebred *Gusb*^{-/-} mice, but no improvement in glenohumeral degeneration. We conclude that *Tlr4*- and/or *C3*- deficiency fail to ameliorate skeletal abnormalities, and other pathways may be involved. RV treatment improves some but not all aspects of bone disease. Radiographs may be an efficient method for future evaluation, as they readily show glenohumeral joint abnormalities.

Keywords

Mucopolysaccharidosis; toll-like receptor; complement; degenerative joint disease; dysostosis multiplex; gene therapy

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1. Introduction

The mucopolysaccharidoses (MPS)¹ are a family of lysosomal storage disorders characterized by deficiencies in enzymes that contribute to the degradation of glycosaminoglycans (GAGs) [1]. The resulting accumulation of GAGs leads to bone and joint disease, respiratory and cardiovascular complications, mental retardation, and hearing and vision impairment. Skeletal abnormalities are referred to as dysostosis multiplex, which results in a limited range of motion, stunted long bones, and difficulty ambulating due to degenerative joint disease (DJD) or loss of joint stability.

MPS VII is a type of MPS due to the deficiency of the enzyme β -glucuronidase (*Gusb*), and will be referred to hereafter as *Gusb*^{-/-}. Skeletal disease is severe in *Gusb*^{-/-} patients [2-6], dogs [6-9], and mice [10-11]. The pathogenesis of skeletal disease is still being investigated, and one hypothesis is that GAGs bind to toll-like receptor 4 (*Tlr4*), resulting in upregulation of inflammatory cytokines such as tumor necrosis factor- α (*Tnf*), interleukin-1 β (*IL1b*), and chemokine (C-C motif) ligand 3 [*Ccl3* or macrophage inflammatory protein 1 α (*MIP1a*)] [12-14]. Indeed, GAGs are structurally similar to the canonical ligand for *Tlr4*, lipopolysaccharide (LPS), and the upregulation of *Tnf*, *IL1b*, and *Ccl3* by GAGs was reduced in microglial cells of *Tlr4*-deficient mice [12]. These inflammatory cytokines are upregulated in blood, synovial fluid and/or cultured fibroblast-like synoviocytes of MPS animals [13-18], and induce expression of destructive proteases [19] such as matrix metalloproteinases (MMPs) and cathepsins that are associated with DJD [20-23]. MPS has also been associated with proliferation of fibroblast-like synoviocytes, upregulation of proteases, and chondrocyte apoptosis in MPS VI rats [13, 16], which likely contribute to the synovial hyperplasia and joint and cartilage degeneration seen in MPS [8-9, 24-25]. The reduced proliferation in the growth plate of *Gusb*^{-/-} mice [26] is likely responsible for the stunting of long bones. It was previously reported that *Gusb*^{-/-} mice that were also deficient in *Tlr4* had improvements in bone lengths and a reduction in synovial *Tnf* RNA levels [14]. However, the effect on degenerative changes in the bones was not assessed.

Another hypothesis for the pathogenesis of disease in MPS is that GAGs activate the complement pathway. We previously demonstrated that complement was activated in the aorta of *Gusb*^{-/-} mice and proposed that the pathogenesis of aortic dilatation might involve complement activation [27], as a variety of carbohydrates can activate complement [28]. Complement is important for immune-complex-induced arthritis [29] and plays a role in *Tlr4* signaling, as deficiency of an inhibitor of complement, *Cd55*, markedly potentiates the effect of LPS on *Tlr4*-dependent cytokine signaling [30-31]. Complement component 3 (C3) is central to the classical, alternative, and lectin pathways, and inhibition of the complement pathway can reduce *Tnf* levels in inflammation [32].

¹Abbreviations: mucopolysaccharidosis (MPS); β -glucuronidase (*Gusb*); glycosaminoglycan (GAG); toll-like receptor 4 (*Tlr4*); complement component 3 (C3); retroviral vector (RV); degenerative joint disease (DJD); tumor necrosis factor- α (*Tnf*); interleukin-1 β (*IL1b*); chemokine (C-C motif); ligand 3 (*Ccl3*); macrophage inflammatory protein 1 α (*MIP1a*); lipopolysaccharide (LPS); matrix metalloproteinase (MMP); hematopoietic stem cell transplantation (HSCT); enzyme replacement therapy (ERT); intravenous (IV); mannose 6-phosphate (M6P); double knock-out (DKO); triple knock-out (TKO); fibroblast growth factor receptor 3 (*Fgfr3*); cathepsin K (CtsK).

Current treatments for some types of MPS patients include hematopoietic stem cell transplantation (HSCT) and/or enzyme replacement therapy (ERT) [33-35]. Neither has prevented the skeletal abnormalities associated with MPS, although ERT has improved ambulation in some MPS I [36], MPS II [37], and MPS VI [38-40] patients. Gene therapy is being tested in animal models [41]. One method involves a neonatal intravenous (IV) injection of a gamma retroviral vector (RV), which transduces liver cells that express the desired enzyme. The enzyme is modified with mannose 6-phosphate (M6P) and secreted into the blood, after which it diffuses to tissues and is taken up by cells via the M6P receptor. Previously, we have observed that a neonatal injection of an RV increased bone lengths and reduced degeneration in *Gusb*^{-/-} dogs [8-9] and increased bone lengths in *Gusb*^{-/-} mice [10]. However, the effect of gene therapy on DJD was not evaluated in *Gusb*^{-/-} mice. The goal of this study was to evaluate the effects of *Tlr4*- and *C3*-deficiency and RV-treatment on the development of skeletal disease in *Gusb*^{-/-} mice.

2. Materials and methods

2.1 Animal care and genotyping

Guidelines set by the National Institutes of Health for the care and use of animals in research were followed. All mice were on a C57Bl/6 background. Genotyping for *Gusb* was done as previously described using a Taqman PCR assay [27] sensitive to the single bp insertion in exon 10 [42]. *C3*^{-/-} mice had a neomycin-resistance gene inserted into the promoter region of *C3* [43] and were generously provided by Drs. Xiabo Wu and John Atkinson of Washington University in St. Louis. Genotyping for *C3* deficiency used a SYBR green mastermix from KAPA Biosystems (Wilmington, MA), and primers that recognized the wild-type *C3* gene (Forward: 5'-TGTTGCCCCAGGTTTGTGA-3' and Reverse: 5'-CCAGGGACTGCCCAAATTT-3') at 61°C, or the *C3* gene with a neomycin insertion (Forward: 5'-CGACAAGACCGGCTTCCA-3' and Reverse: 5'-AAGCGAAACATCGCATCGA-3') at 61°C. *Tlr4*^{-/-} mice had a 74 kb deletion that included the *Tlr4* coding sequence [44], and were obtained from Jackson labs (Bar Harbor, ME; B6.B10ScN-*Tlr4*^{lps-del}/JthJ; stock number #007227). Genotyping for *Tlr4* used SYBR green primers that recognized the wild-type *Tlr4* gene (Forward: 5'-AGAAATTCCTGCAGTGGGTCA-3' Reverse: 5'-TCTCTACAGGTGTTGCACATGTCA-3') at 61°C, or the *Tlr4* mutation (Forward: 5'-GCAAGTTTCTATATGCATTCTC-3' and Reverse: 5'-CCTCCATTTCCAATAGGTAG-3') at 63°C. Some *Gusb*^{-/-} mice were injected IV at 2-3 days after birth with 1×10¹⁰ transducing units/kg of the RV designated hAAT- cGusb-WPRE that expresses canine *Gusb* and contains the human α₁-antitrypsin promoter [45], which allowed them to survive and breed. Some RV-treated *Gusb*^{-/-} mice were bred with *C3*^{-/-} or with *Tlr4*^{-/-} mice to generate obligate heterozygotes, which were then crossed to generate *Gusb*^{-/-} *Tlr4*^{-/-}, *Gusb*^{-/-} *C3*^{-/-}, or *Gusb*^{-/-} *Tlr4*^{-/-} *C3*^{-/-} mice.

2.2 Radiographs

At the time of sacrifice, mice were anaesthetized by injection of ketamine/zylazine as reported previously [27], perfused with 20 ml of PBS, and died of exsanguination. Bones were frozen at -20°C, and radiographs were obtained and scanned into the computer. Femur

and tibia measurements were obtained as shown in Supplemental Fig. 1A-C. Arm radiographs were blinded as to the genotype, and the glenohumeral joint space was measured as shown in Supplemental Fig. 1D and scored for dysplasia and irregularities as shown in Supplemental Fig. 2. For dysplasia of the proximal humerus, 0 represents a perfectly spherical ball shape; 1 represents a slightly flattened sphere but the ball shape was still apparent; 2 represents an oval shape; and 3 represents a shallow oval with an almost completely flat surface. For glenoid cavity dysplasia, 0 represents a flattening of 0-4% of the surface of the glenoid cavity; 1 represents a 5-29% flattening; 2 represents a 30-59% flattening; and 3 represents a 60-100% flattening. Irregularities of both the proximal humerus and glenoid cavity were scored with the following criteria: 0 represents irregularity of 0-10% of the subchondral surface; 1 represents 11-29% irregularity of the surface; 2 represents 30-60% irregularity; and 3 represents more than 60% irregularity.

2.3 Histochemistry

Bones were fixed for 10-14 days in a solution of phosphate buffered saline with 10% formalin and decalcified for ~24 hours in Formical-2000 (Decal-Bone, Talman, NY) until the bones were pliable. Tissues were embedded in paraffin and 6- μ m sections were stained with Masson's Trichrome. Slides were photographed with an Olympus Nanozoomer 2.0-HT system and NDP imaging software.

2.4 Statistical Analysis

Significance for values that were continuous (leg measurements and joint space) was assessed using One-Way ANOVA with Holm-Sidak post-hoc analysis using SigmaPlot 12.0 (Sigma-Aldrich, St. Louis, MO). Significance for values that were non-continuous (glenohumeral DJD scores) was compared using ANOVA on ranks with Dunn's post-hoc analysis.

3. Results

Gusb^{-/-} mice were bred with *Tlr4*^{-/-} mice and *C3*^{-/-} mice to generate doubly and triply-deficient mice. *Gusb*^{+/-} *Tlr4*^{+/+} *C3*^{+/+} mice will be referred to hereafter as normal mice, as previous studies have shown that mice that are heterozygous for *Gusb* are phenotypically normal [46]. *Gusb*^{-/-} *Tlr4*^{+/+} *C3*^{+/+} mice will be referred to hereafter as purebred *Gusb*^{-/-} mice. *Gusb*^{-/-} mice that were also deficient in *Tlr4* but were normal for the *C3* locus (*Gusb*^{-/-} *Tlr4*^{-/-} *C3*^{+/+}) will be referred to hereafter as *Gusb-Tlr4* DKO (double knock-out) mice, although neither of these mutant strains were actually generated with homologous recombination. *Gusb*^{-/-} *Tlr4*^{+/+} *C3*^{-/-} mice will be referred to as *Gusb-C3* DKO mice. *Gusb*^{-/-} *Tlr4*^{-/-} *C3*^{-/-} mice will be referred to as *Gusb-Tlr4-C3* TKO (triple knock-out) mice.

3.1 Radiographs of the femur and tibia

Representative radiographs of the femurs at 3 months of age are shown in Fig. 1A, where mice of all *Gusb*^{-/-} genotypes had significantly shorter femurs than normal mice. Fig. 1B shows that purebred *Gusb*^{-/-} mice had femurs that were only 77 \pm 1% as long as purebred normal mice ($p < 0.001$). *Gusb-Tlr4* DKO, *Gusb-C3* DKO, and *Gusb-Tlr4-C3* TKO mice had no improvement in lengths compared with purebred *Gusb*^{-/-} mice, and bones remained

statistically shorter than in purebred normal mice ($p < 0.001$). Similarly, purebred *Gusb*^{-/-} tibias were $84 \pm 5\%$ the length of normal tibias ($p < 0.001$), and there was no improvement in the lengths of *Gusb-Tlr4* DKO, *Gusb-C3* DKO, and *Gusb-Tlr4-C3* TKO mice compared to purebred *Gusb*^{-/-} mice. Purebred *Gusb*^{-/-} mice had femurs that were $124 \pm 5\%$ as wide as normal mice ($p < 0.001$) (Fig. 1D), and the femurs of *Gusb-Tlr4* DKO, *Gusb-C3* DKO, and *Gusb-Tlr4-C3* TKO mice remained wide ($p < 0.001$ vs. normal). *Gusb*^{-/-} mice that were heterozygous for *Tlr4* and/or *C3* had values similar to those of purebred *Gusb*^{-/-} mice (data not shown). Similarly, normal *Gusb*^{+/-} mice that were +/- or -/- for *Tlr4* and/or *C3* had similar values to purebred *Gusb*^{+/-} mice (data not shown). Degeneration of the stifle joint was difficult to observe on radiographs in purebred *Gusb*^{-/-} mice compared with normal mice (data not shown).

3.2 Histochemistry of the stifle joint

Histochemistry was evaluated to determine bone and cartilage quality in the stifle joint. Normal mice (Fig. 2A-B) displayed a strong line of solid subchondral bone, with normal-sized chondrocytes in the articular cartilage and growth plate. Purebred *Gusb*^{-/-} mice (Fig. 2C-D) displayed thinning of the subchondral bone in both the distal femur and proximal tibia, and the articular cartilage and growth plate were thickened due to GAG accumulation. *Gusb-Tlr4* DKO, *Gusb-C3* DKO, and *Gusb-Tlr4-C3* TKO mice (Fig. 2E-J) exhibited similar bone thinning and thickening of articular cartilage and the growth plate as did purebred *Gusb*^{-/-} mice.

3.3 Radiographs of the glenohumeral joint

3.3.1 Joint space measurements—Radiographs of mouse arms displayed shortened humeri and ulna/radii lengths in all *Gusb*^{-/-} genotypes compared to normal mice (data not shown). Representative examples of radiographs of the glenohumeral joint are shown in Fig. 3A to 3E, and joint space measurements were obtained to determine the distance between the bones of the proximal humeri and the bones of the glenoid cavity as shown in Supplemental Fig. 1D, and the average values for several animals are shown in Fig. 3F. Normal mice had an average joint space of 0.10 ± 0.03 mm, which was significantly narrower than for purebred *Gusb*^{-/-} mice with an average joint space of 0.46 ± 0.06 mm ($p < 0.001$). The joint space of *Gusb-Tlr4* DKO, *Gusb-C3* DKO, and *Gusb-Tlr4-C3* TKO mice was not significantly improved compared to that of purebred *Gusb*^{-/-} mice.

3.3.2 Dysplasia—Dysplasia of the proximal humerus and glenoid cavity was scored as flattening of the articular surfaces from 0 (normal) to +3 (severely abnormal) as stated in the methods and shown in Supplemental Fig. 2. For the proximal humerus, normal mice scored an average of 0.0 ± 0.0 . Purebred *Gusb*^{-/-} mice scored significantly higher than normal mice with an average of 2.0 ± 0.8 ($p < 0.05$), and there was no significant improvement in *Gusb/Tlr4* DKOs (2.4 ± 0.7), *Gusb/C3* DKOs (2.5 ± 0.7), or *Gusb/Tlr4/C3* TKOs (1.9 ± 0.6) compared with purebred *Gusb*^{-/-} mice ($p > 0.05$). For dysplasia of the glenoid cavity, normal mice scored an average of 0.0 ± 0.0 , which was significantly lower than for purebred *Gusb*^{-/-} mice (1.5 ± 1.2 ; $p < 0.05$). Again, no significant improvement was seen in *Gusb/Tlr4* DKO (1.9 ± 0.9), *Gusb/C3* DKO (2.3 ± 0.8), or *Gusb/Tlr4/C3* TKO (2.2 ± 0.9) mice compared to purebred *Gusb*^{-/-} mice ($p > 0.05$).

3.3.3 Surface irregularities—Irregularities of the subchondral bone are another manifestation of MPS that appear on radiographs as a lack of a solid, traceable edge or osteophyte formation, and can be due to inadequate bone formation, erosions, or new bone formation. Irregularities were scored as stated in the methods and as shown in Supplemental Fig. 2. For both the proximal humerus and the glenoid cavity, normal mice scored an average of 0.3 ± 0.3 . Purebred *Gusb*^{-/-} mice scored significantly higher in both areas with an average of 1.8 ± 1.0 for the humerus ($p < 0.05$ vs. normal), and 2.5 ± 1.8 for the glenoid cavity ($p < 0.01$ vs. normal). No significant improvement was seen in either area between *Gusb/Tlr4* DKO, *Gusb/C3* DKO, or *Gusb/Tlr4/C3* TKO mice and purebred *Gusb*^{-/-} mice.

3.4 Histochemistry of the humerus and glenoid cavity

3.4.1 Articular cartilage and bone quality—As was observed in the stifle joint, normal mice (Fig. 4A-B) exhibit a strong rim of bone underlying the articular cartilage, with no excess GAGs in either the articular cartilage or the growth plate. In contrast, purebred *Gusb*^{-/-} mice (Fig. 4C-D) lack a solid rim of subchondral bone, and display thickened articular cartilage and growth plate due to the accumulation of GAGs. No improvement was seen in *Gusb-Tlr4* DKO, *Gusb-C3* DKO, and *Gusb-Tlr4-C3* TKO mice (Fig. 4E-J), as they similarly lacked the strong line of bone in normal mice, and displayed thickened cartilage.

3.4.2 Synovial hyperplasia—In addition to differences in the bone and cartilage, differences in the synovium can also be observed between normal and purebred *Gusb*^{-/-} mice, as shown in Supplemental Fig. 3. The synovium of normal mice is a few cell layers thick, while the hyperplastic synovium in purebred *Gusb*^{-/-} mice is several cell layers thick, especially in the condylar neck region of the humerus. The synovium appears similarly hyperplastic in *Gusb-Tlr4* DKO, *Gusb-C3* DKO, and *Gusb-Tlr4-C3* TKO mice, although not all mice had a region on the slide that contained synovium, making this difficult to score.

3.5 Effect of RV-treatment on bones

During the breeding process, some *Gusb*^{-/-} mice that were +/+ or +/- for *Tlr4* and *C3* were treated with a neonatal IV injection of RV. The average serum *Gusb* activity of these RV-treated mice was 1191 ± 1201 Units/ml, which is 85-fold the value in heterozygous normal mice (Fig. 5A). Radiographs of these RV-treated mice at an average age of 8.4 months of age were compared with those of purebred *Gusb*^{-/-} and *Gusb*^{+/-} normal mice at 6 months of age for bone lengths and joint degeneration. Representative radiographs of the femur and evaluation of bone measurements are shown in Fig. 5B-E. Purebred *Gusb*^{-/-} mice had femurs that were $124 \pm 6\%$ the width of normal femurs ($p < 0.001$). RV-treated mice had significantly narrower femurs than purebred *Gusb*^{-/-} mice ($p < 0.001$) and were $103 \pm 4\%$ of normal ($p = 0.324$). Purebred *Gusb*^{-/-} mice had femurs and tibias that were $76 \pm 1\%$ and $81 \pm 3\%$, respectively, the length of normal femurs and tibias ($p < 0.001$ vs. normal for both). Although RV-treated mice had significantly longer femurs and tibias at $84 \pm 4\%$ and $89 \pm 2\%$ of normal, respectively, ($p < 0.001$ for both vs. untreated purebred *Gusb*^{-/-} mice), they were significantly shorter than those of normal mice ($p < 0.001$ for both). Degeneration in the stifle joint could not be clearly seen on radiographs, so was not evaluated.

The arms of RV-treated mice were longer than those of purebred *Gusb*^{-/-} mice, but were not as long as normal mice (data not shown), as has been previously reported [10]. The joint space was measured as shown in Supplemental Fig. 1D, and as shown in Fig. 5F and 5G. Purebred *Gusb*^{-/-} mice had an average joint space of 0.71±0.18 mm, which was significantly wider than that of normal mice with 0.17±0.10 mm ($p < 0.001$ vs. *Gusb*^{-/-}) and that of RV-treated mice with 0.40±0.18 mm ($p < 0.001$ vs. *Gusb*^{-/-}; $p < 0.001$ vs. normal). Radiographs of the glenohumeral joint were scored on a scale from 0 (normal) to +3 (severely abnormal) for dysplasia and surface irregularities, and values are shown in Fig. 5H. Purebred *Gusb*^{-/-} mice scored significantly higher than normal mice in dysplasia and surface irregularities of the humerus and glenoid cavity ($p < 0.01$). RV-treated mice scored significantly higher than normal mice in every aspect that was scored ($p < 0.01$). The only significance found between RV-treated and purebred *Gusb*^{-/-} mice was in irregularities of the humerus, where RV-treated mice scored more severely at 1.7±0.9 compared to purebred *Gusb*^{-/-} mice at 1.3±0.8 ($p < 0.05$). This may relate to the fact that RV-treated *Gusb*^{-/-} mice were older than untreated mice, and thus had more time to develop degenerative changes.

4. Discussion

This study was conducted to evaluate the effects of *Tlr4*- and *C3*-deficiency, and RV-treatment on skeletal abnormalities in *Gusb*^{-/-} mice. Skeletal disease is still prevalent in MPS patients after HSCT or ERT, which greatly reduces their quality of life. The failure to correct skeletal disease is likely due to the inability of blood-derived cells or enzyme to enter avascular regions such as cartilage. A hypothesis of this project is that GAGs bind to *Tlr4* and/or activate the complement pathway, resulting in up-regulation of cytokines that induce the expression of proteases that contribute to DJD. Confirmation of the role of these pathways in MPS joint disease could lead to their inhibition through drugs as a possible treatment.

4.1. Deficiency of *Tlr4* and/or *C3* does not prevent bone and joint disease in MPS VII mice

To test the role of *Tlr4* and *C3* in the pathogenesis of joint disease, *Gusb*^{-/-} mice were bred with mice that were deficient in *Tlr4* and *C3* to generate doubly and triply deficient mice, and the effects on bone disease were determined. In this study, *Gusb-Tlr4* DKO, *Gusb-C3* DKO, and *Gusb-Tlr4-C3* TKO mice showed no significant improvement in lengths of the long bones of the arms and legs, in width of the femur, or in scores of dysplasia and bone irregularities of the glenohumeral joint when compared with purebred *Gusb*^{-/-} mice.

The failure to prevent bone shortening in *Gusb-Tlr4* DKO mice in this study is in contrast to the study of Simonaro *et al.* [14], which reported that the lengths of femurs and tibias in male *Gusb-Tlr4* DKO mice were approximately 1.3-fold that in purebred *Gusb*^{-/-} mice, and approached the values found in *Gusb*^{+/+} *Tlr*^{-/-} mice. Both studies used *Gusb*^{-/-} mice derived from Dr. Mark Sands at Washington University, and *Tlr4*^{-/-} mice of the strain indicated in the Methods section from Jackson Laboratories (Calogera Simonaro, personal communication). It is unlikely that inaccurate genotyping on our part is responsible for this difference, as some *Gusb-Tlr4* DKO mice that were bred and genotyped independently by Dr. William Sly using a different PCR technique but were radiographed by our laboratory had similar bone lengths and DJD scores as the animals bred, genotyped, and radiographed

in our laboratory (data not shown). Although we pooled results for male and female mice in this study, our groups had a similar percentage of mice of each gender, and we failed to observe significant differences between male and female mice or between groups when animals of the same gender were analyzed separately (data not shown). Our study did evaluate mice that were older (3 to 6 months) than in the study by Simonaro *et al.* (1.5 months), so it is possible that the age difference was responsible for the discrepancy. It is also possible that genetic drift occurred in either the *Gusb*^{-/-} or the *Tlr4*^{-/-} colonies from the time when each was obtained from their source, which could contribute to the different results of the two studies. Indeed, the *Gusb*^{-/-} mice in our study had femur lengths of 1.2 cm at 3 months, while the *Gusb*^{-/-} mice in the previous study had femur lengths of ~1 cm at 6 weeks [14], suggesting that differences could have been due to age or genetic background.

We conclude that *Tlr4*-deficiency does not protect *Gusb*^{-/-} mice from stunted bones or DJD, and that inhibition of *Tlr4* would not be effective against skeletal disease should such drugs be developed in the future. Indeed, the work of Ausseil *et al.* [12] demonstrated that although *Ccl3* expression in response to GAGs in cultured microglial cells was only ~20% as high in *Tlr4*^{-/-} cells as in normal cells, *Ccl3* expression in *Tlr4*^{-/-} cells that were stimulated with GAGs remained ~20-fold that of cells that were not stimulated with GAGs, suggesting that the *Tlr4* pathway was not the only GAG-activated pathway. Furthermore, although *Ccl3* mRNA in brain was reduced at 3 months or earlier in MPS IIIB mice that were deficient in *Tlr4* compared to MPS IIIB mice normal for *Tlr4*, that difference was no longer present at 8 months of age, again suggesting that *Tlr4* is not the only GAG-activated pathway that is inducing cytokines.

Thus, although GAGs can clearly induce cytokine expression via the *Tlr4* pathway, they can also induce cytokines independently of *Tlr4*, which may explain why *Tlr4*-deficiency did not ameliorate bone disease in *Gusb*^{-/-} mice in our study. *Tnf* is an important mediator of *Tlr4* signaling, and inhibition of *Tnf* with an inhibitory antibody can reduce bone disease in MPS VI rats [14-15]. It would have been informative to test *Gusb*-*Tlr4* DKO or *Gusb*-*Tlr4*-*C3* TKO mice for *Tnf* RNA or protein levels in the synovium, bone, and cartilage, as *Tnf* can be induced by other signaling pathways. However, isolation of these tissues from mice is problematic due to their small size and was not attempted in this study.

Complement is another pathway that can be activated by GAGs [28] and is activated in the aorta of *Gusb*^{-/-} mice [27], and C3 is critical component of the classical, alternative, and lectin pathways. However, deficiency of C3 alone or combined deficiency of *Tlr4* and C3 did not ameliorate bone and joint disease in this study. We had previously postulated that GAGs might activate fibroblast growth factor receptor 3 (*Fgfr3*), as GAGs are important in the interaction of *Fgfr3* with its ligand, and mutations that increase activation of *Fgfr3* result in achondroplastic dwarfism. However, deficiency of *Fgfr3* also did not prevent stunting of bones in *Gusb*^{-/-} mice [26]. It is possible that GAGs are activating still another signaling pathway, which explains why deficiency of *Tlr4*, *C3*, or *Fgfr3* did not prevent skeletal manifestations of disease.

4.2. Possible role of downstream mediators in bone and joint disease

An alternative explanation is that GAGs can cause bone and joint disease via a more direct mechanism such as activation of cathepsin K (CtsK), which is a protease that degrades bone and cartilage. Since CtsK is activated directly by chondroitin sulfate [47], a GAG that accumulates in MPS VII, it may be activated independently of signaling pathways. Indeed, CtsK mRNA and/or enzyme activity were elevated in the annulus fibrosus of the spine of MPS VII dogs [6] and in the aorta of MPS VII mice [27]. Our future studies will test if bone disease can be reduced in MPS VII mice by the CtsK-specific inhibitor odanacitib, which is currently in clinical trials for treating osteoporosis [48]. Interestingly, pentosan polysulfate, a drug that promotes chondrogenesis, was recently reported to reduce bone disease in MPS VI rats [17-18]. Although the authors demonstrated a reduction in serum cytokines and hypothesized that its mechanism reduced inflammation, it may be interesting to test if pentosan polysaccharide directly affects CtsK. Deficiency of other proteases such as cathepsin S or MMP12 did not ameliorate bone disease in *Gusb*^{-/-} mice [27].

4.3. RV-treatment

Gusb^{-/-} mice that were treated neonatally with an IV injection of an RV expressing canine *Gusb* were evaluated to determine the effects on bone disease. Bone lengths were partially improved relative to untreated *Gusb*^{-/-} mice, which was consistent with our previous results in mice treated with the same vector that had similar levels of expression [10] and with our results in RV-treated *Gusb*^{-/-} dogs [8-9]. In contrast, manifestations of DJD such as dysplasia and bone irregularities were not prevented in the RV-treated mice. Similarly, osteophyte formation was not prevented at any sites that was evaluated and dysplasia still occurred in the proximal femur, proximal tibia, and cervical spine of RV-treated *Gusb*^{-/-} dogs, although dysplasia was reduced in the acetabulum and distal femur [8-9]. Subchondral bone irregularities were improved at 1 year but not at 8 years in RV-treated *Gusb*^{-/-} dogs [8-9]. Thus, this study shows that evaluation of *Gusb*^{-/-} mice for DJD using plain radiographs of the glenohumeral joint was quite informative, and demonstrated results in RV-treated *Gusb*^{-/-} mice that were largely consistent with those in the RV-treated *Gusb*^{-/-} dogs. Similar radiographic evaluation may therefore be useful for future studies that test the efficacy of a particular treatment on DJD in mice with MPS.

The failure to prevent most manifestations of DJD after neonatal gene therapy with an RV in *Gusb*^{-/-} mice leads to our prediction that ERT will be similarly limited in its efficacy, even if started in the neonatal period. Although *Gusb* enzyme can diffuse to the synovium of *Gusb*^{-/-} dogs [9] and to the edge of the cortex of bone of *Gusb*^{-/-} mice [10] after neonatal gene therapy with an RV, articular cartilage of RV-treated *Gusb*^{-/-} dogs did not have *Gusb* activity detected with a histochemical stain (E. Xing, K. Ponder, unpublished data). Low levels of enzyme in cartilage in RV-treated dogs may be due to the low levels of *Gusb* activity in synovial fluid at 7±5 U/ml (2±2% of the level in blood) [9] and the dense nature of cartilage that reduces diffusion, as well as the lack of a blood supply.

4.3. Future implications

This study suggests that neither Tlr4 nor C3 is essential for the pathogenesis of bone and joint disease in MPS VII mice, and that inhibition of these signaling pathways will not

prevent stunted bones or DJD in patients. Neonatal IV injection of an RV can partially improve bone lengths, but has no effect on DJD. Our future studies will inject vectors directly into joints of MPS dogs and monitor the effects on skeletal disease, as intra-articular injection of enzyme has reduced lysosomal storage in MPS I dogs [49].

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Research highlights

1. Deficiency of Tlr4 or Complement C3 does not ameliorate bone disease in MPS VII mice.
2. Neonatal gene therapy with an IV injection of a retroviral vector improves bone lengths but does not prevent degenerative changes in MPS VII mice.

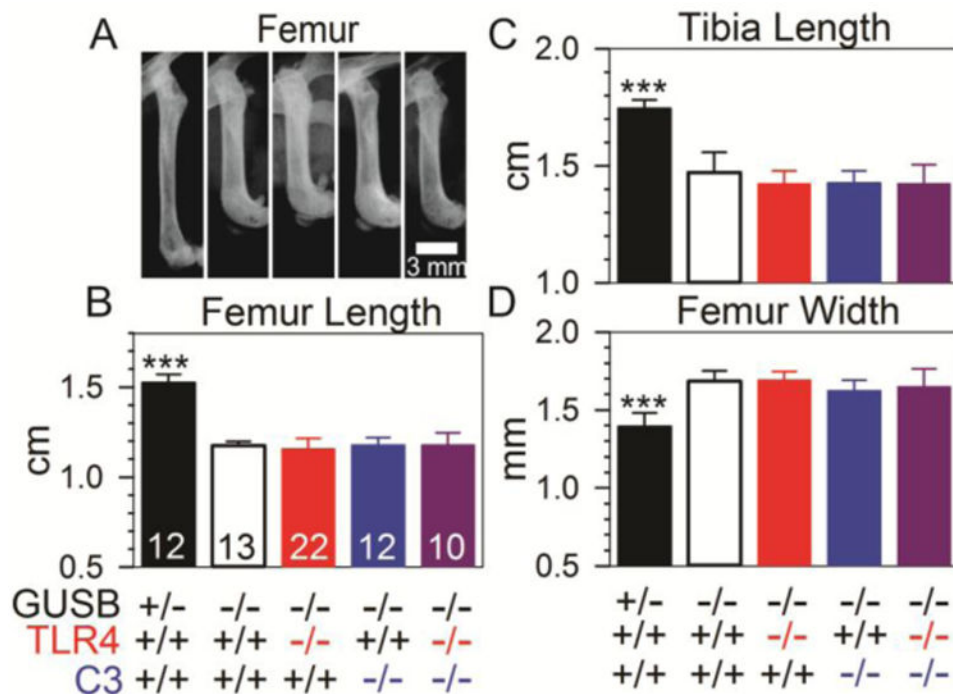


Fig. 1. Radiographical evaluation of the femur and tibia

Radiographs of the legs were obtained. **A.** Representative radiographs of the femur of male mice of the indicated genotype at 3 months of age are shown. The proximal femur is at the top. **B-D.** Measurements of femur length, tibia length, and femur width at 3-6 months of age are shown for the indicated number of mice, which included both males and females. Statistical comparison between purebred *Gusb*^{-/-} mice and the other groups used ANOVA with Holm-Sidak post-hoc analysis, and *** signifies a p-value <0.001.

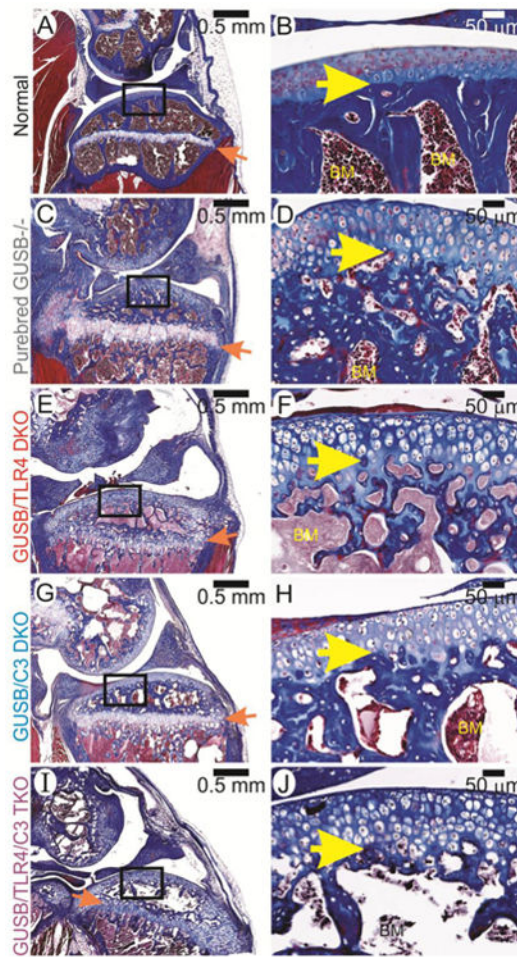


Fig 2. Histochemistry of the femur and tibia

Sections of stifle joints from male mice at 3 months of age were stained with Masson's Trichrome as stated in the methods. Representative photographs were taken at low (left) and high (right) power. The groups are indicated on the far left. In the left column, the distal femur is at the top while the proximal tibia is at the bottom. The orange arrows indicate the growth plate. The boxes outline the area magnified in the right column, in which the bone marrow (BM) is indicated, and the yellow arrows point out the junction between articular cartilage and bone.

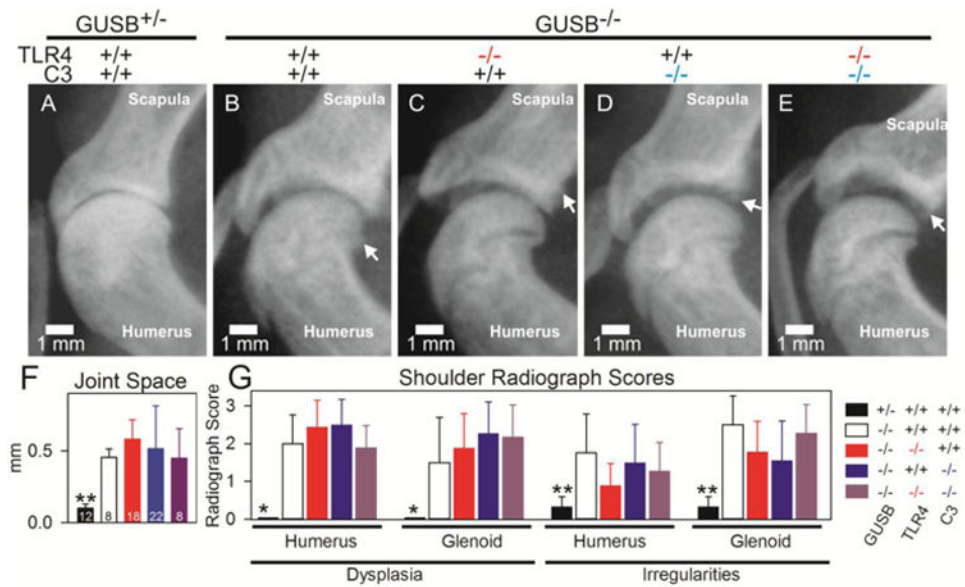


Fig 3. Radiographical evaluation of the humerus and glenoid cavity

Radiographs of the arms were obtained at 3 months of age. **A-E**. Representative radiographs of the glenohumeral joint of male mice of the indicated genotype are shown. The joint space appears as the black region between the glenoid cavity of the scapula and the humerus. Arrows indicate irregular regions of the subchondral bone. **F**. Joint space measurements were performed for the indicated number of mice, which included both males and females. Statistical comparisons between purebred *Gusb*^{-/-} mice and the other groups were done using One-way ANOVA with Holm Sidak's post-hoc analysis. * signifies a p-value between 0.01 and 0.05, ** signifies a p-value of 0.001 to 0.01, and *** represents a p value < 0.001. **G**. Dysplasia and irregularities of the subchondral bone were scored for the same mice as in panel F for the humerus and glenoid cavity as detailed in the Methods section and illustrated in Supplemental Fig. 2. A score of 0 is normal and +3 is severely abnormal. Statistical comparisons were done using Kruskal-Wallis ANOVA on ranks with Dunn's post-hoc analysis.

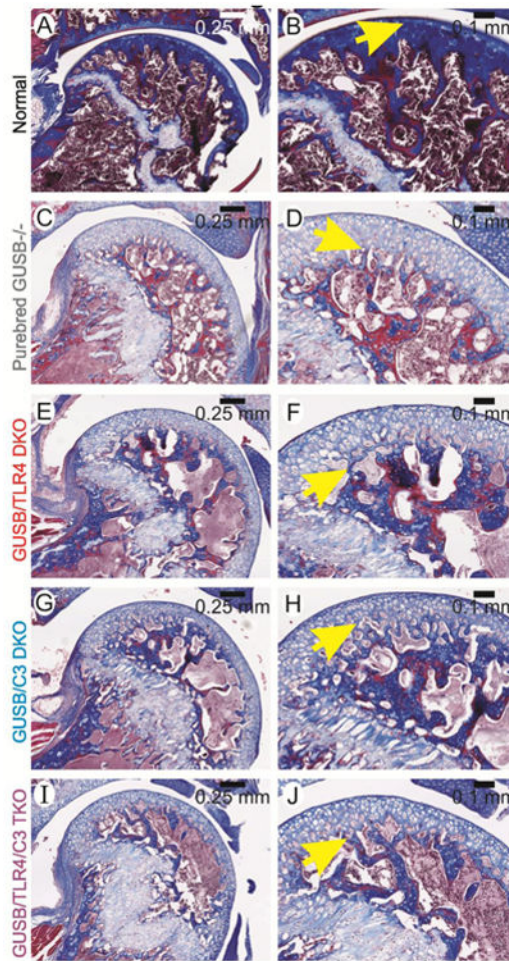


Fig 4. Histochemistry of the humerus and glenoid cavity

Glenohumeral joints from mice at 3 months of age were stained with Masson's Trichrome. Genotypes are indicated on the far left. Representative photographs were taken at low (left) and high (right) power. In the left column, the proximal humerus is shown, and is magnified in the right column panels, where the yellow arrows indicate the junction between articular cartilage and bone.

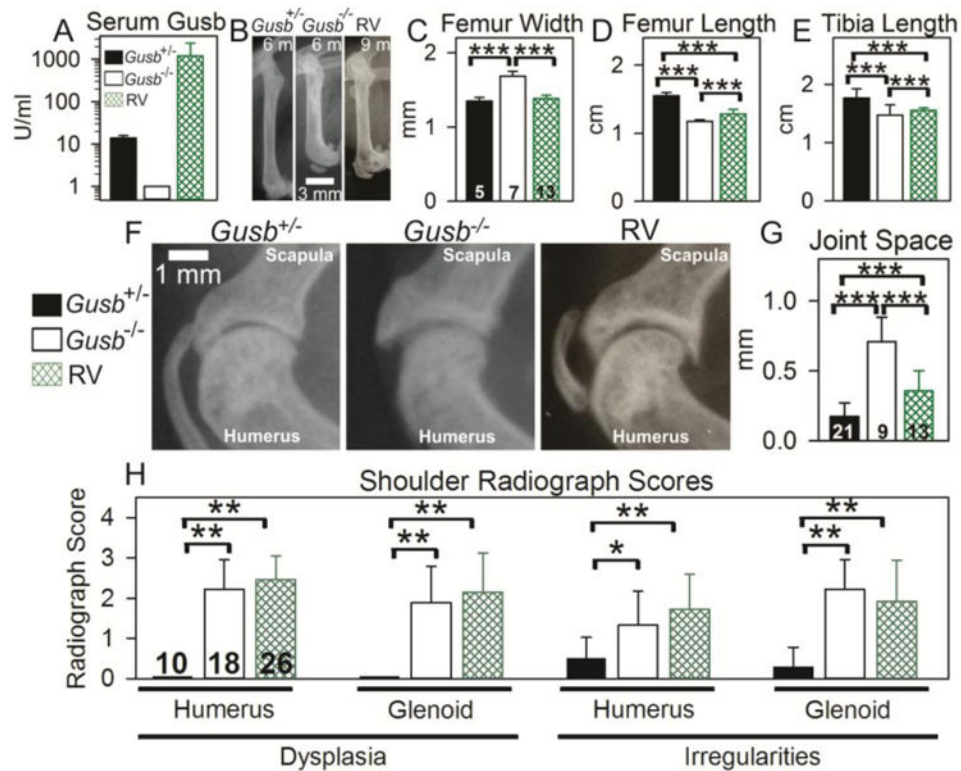


Fig. 5. Radiographical evaluation of RV-treated mice

Some *Gusb*^{-/-} mice were injected with RV at 2 to 3 days after birth (RV), and bones were evaluated at an average age of 8.4 months. These mice were either +/- or +/+ for *Tlr4* and *C3*. Other *Gusb*^{-/-} and *Gusb*^{+/-} mice were untreated and bones were evaluated at the younger age of 6 months. **A. Serum Gusb activity.** Average Gusb serum activity was tested at 2 months of age. **B.** Representative radiographs of femurs with the groups indicated above the radiographs. The proximal femur is at the top and the age is shown in months (m). **C-E.** Femur width and length and tibia length were measured for the indicated number of mice. **F.** Representative radiographs of the glenohumeral joint are shown. **G.** Joint space was measured for the indicated number of mice. **H.** Dysplasia and surface irregularities were scored as stated in the methods and illustrated in Supplemental Fig.2, where 0 indicates normal and 3 indicates severely abnormal. The indicated number of mice were evaluated. Statistical comparison for panels C-E and panel G used one-way ANOVA with Holm-Sidak post-hoc analysis, and *** signifies a p-value <0.001. Statistical comparisons for panel H used Kruskal-Wallis ANOVA on ranks with Dunn's post-hoc analysis, and * indicates a p value of 0.01 to 0.05 and ** indicates p<0.01.