Excessive growth hormone expression in bGH transgenic mice adversely alters bone architecture and quality

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Introduction

Growth hormone (GH) is one of the important anabolic hormones involved in regulation of longitudinal bone growth¹ by causing chondrocyte proliferation at the epiphyseal cartilage in the growth plate region. Acromegaly is a disease with abnormal skeletal tissue and organ growth². It is caused by excessive GH expression after puberty². Acromegaly have higher prevalence of patients vertebral fractures despite having normal bone mineral density (BMD), suggesting that BMD alone in not a sufficient indicator of fracture risk. The cause of this high incidence of fracture remains ill-defined.

Results



Aims

- 1. To evaluate the skeletal changes in giant bovine GH overexpressing transgenic mice (bGH)
- 2. To establish the usefulness of bGH mice as a mouse model for the study of skeletal changes in acromegaly.

Methods

Micro-CT analysis of tibia of bGH and Wild Type (WT) mice:

5-month-old male bGH and WT mice (n=7/group) were provided by Prof. John Kopchick (Ohio University, USA). Tibia were scanned with high resolution (5µm pixel size) micro-CT. Bone histomorphometric analyses were performed by SkyScan software (CT-Analyzer v.1.5.1.3). Trabecular bone parameters determined included bone volume (BV)/tissue volume (TV) percentage (BV/TV), %), trabecular thickness (Tb. Th), trabecular bone pattern factor (TBPf). Cortical bone parameters determined included tissue perimeter (T. Pm), cross-sectional thickness (Cs. Th) and medullary area.

Fig 1. Weights of bGH and WT mice. Values are mean \pm SD of n=7 mice/group.***p<0.001.

Fig 2. Lengths of bones (tibia, femur and vertebra) measured using micro-CT. Values are mean ± SD of n=7 mice/group. ***p<0.001

Bone architecture of bGH transgenic mice



Fig 3. Trabecular and cortical microarchitecture in WT and bGH transgenic mice. (A,B,C,D) BV/TV (A), Tb.Th (B), Tb.N (C) and Tb. Pf (D) were measured in the proximal metaphysis of WT and bGH mice (E,F,G) T. Pm (E), Ct. Th (F) and medullary area (G) were measured in the mid-diaphysis cortical bone of WT and bGH mice

Bars represent mean ± SD of n=7 mice/group. *p<0.05, ***p<0.001

Measurement of BMD:

BMD analysis was performed in lumbar vertebrae (L4) using SkyScan software.

Mechanical testing:

Three-point bending tests on femora of WT and bGH mice were performed by as previously described³. Calculations of bone mechanical properties including Young's modulus and ultimate stress.

Histological analysis osteoclast Of activity:

Tibia from bGH and WT mice were fixed in 4% formaldehyde and decalcified. 8µm-thick sections were stained for tartrate-resistant acid phosphatase (TRAP), a marker of osteoclast activity and counterstained with haematoxylin solution. TRAP staining was quantified using ImageJ software. Osteoclast activity was expressed as % TRAP positive stain.

- 1. The decreases in BV/TV, trabecular number and cortical thickness in bGH mice compared to WT mice indicate that bGH mice have low bone mass.
- 2. The significant increase in trabecular pattern factor indicates less inter-trabecular connectivity, suggesting deterioration of trabecular bone in bGH mice.



Statistical analysis:

Statistical analyses were performed using unpaired (two-tailed) t test with GraphPad software, using p<0.05 as significant.

References

Conclusion

1. Ohlsson C et al. JEndocrinol. 1992; 133(2):291-300 2. Melmed S. J Clin Invest. 2009; 119(11):3189-202 3. Marenzana M et al. Arthritis Rheum. 2011; 63(8):2385-95.

High serum GH levels negatively affect bone architecture and quality. bGH transgenic mice are a useful model to understand the mechanisms involved in the skeletal changes observed in acromegaly patients.