

# *PCMD MicroCT Imaging Core Learning Lunch Series*

## MicroCT 101: How to get most out of your scans

X. Sherry Liu, Ph.D  
Wei-Ju Tseng, M.S

January 17<sup>th</sup>, 2018

Mckay Orthopaedic Research Laboratory  
Department of Orthopaedic Surgery  
Perelman School of Medicine  
University of Pennsylvania  
Philadelphia, PA



Penn

McKay Orthopaedic Research Laboratory



# What is $\mu$ CT?

- A non-destructive technique to produce 3D images of very high resolution using X-ray imaging and computed tomography

## Recommended Reading:

Guidelines for assessment of bone microstructure in rodents using micro-computed tomography

Bouxsein ML, Boyd SK, Christiansen BA, Guldborg RE, Jepsen KJ, Müller R.

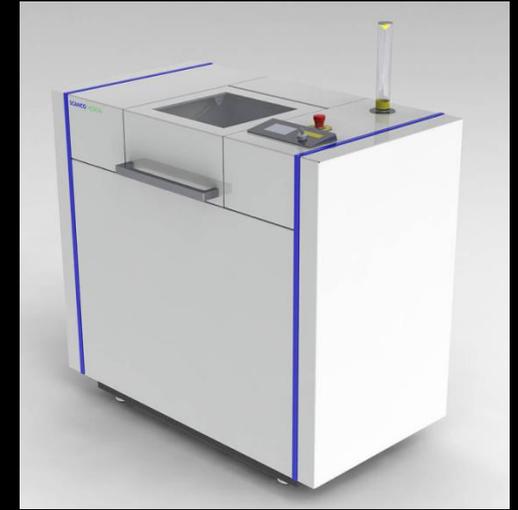
J Bone Miner Res. 2010 Jul;25(7):1468-86. doi: 10.1002/jbmr.141.

- Disease progression
- Drug treatments
- Input to micro finite element ( $\mu$ FE) models to estimate the mechanical properties of bone van Rietbergen+1998, Schulte+2011



# Available Scanners at PCMD $\mu$ CT Imaging Core

- Specimen  $\mu$ CT
  - $\mu$ CT 35
  - $\mu$ CT 50
- *In Vivo*  $\mu$ CT
  - vivaCT 40
  - vivaCT 75
- Clinical  $\mu$ CT
  - XtremeCT II



# PCMD $\mu$ CT Scanners

	$\mu$ CT 35	$\mu$ CT 50	vivaCT 40	vivaCT 75	XtremeCT II
Use	Specimen	Specimen	In Vivo Rodent	In Vivo Rodent	Clinical
X-Ray Source	30 - 70 kVp	30 - 90 kVp	30 - 70 kVp	30 - 70 kVp	68 kVp
Max Scan Size	37.9 x 120 mm ( $\emptyset$ x L)	50 x 120 mm ( $\emptyset$ x L)	38.9 x 145 mm ( $\emptyset$ x L)	79.9 x 145 mm ( $\emptyset$ x L)	140 x 200 mm ( $\emptyset$ x L)
Max Specimen Size	75.8 x 140 mm ( $\emptyset$ x L)	100 x 160 mm ( $\emptyset$ x L)	80 x 500 mm ( $\emptyset$ x L)	80 x 500 mm ( $\emptyset$ x L)	170 mm ( $\emptyset$ )
Best image voxel size	3.5 $\mu$ m ( $\emptyset$ :7 mm)	1.5 $\mu$ m ( $\emptyset$ : 3 mm)	10.5 $\mu$ m ( $\emptyset$ :21.5 mm)	19 $\mu$ m ( $\emptyset$ : 38.9 mm)	60 $\mu$ m
Location	McKay Lab	VA Hospital	McKay Lab	VA Hospital	CHOP CTRC nutrition and growth lab

# Next generation *in vivo* $\mu$ CT scanner

- Scanco vivaCT 80
  - Location: McKay Lab
  - *In vivo*  $\mu$ CT scanner
  - Best image voxel size:
    - 10.5  $\mu$ m isotropic voxel size @  $\emptyset$  21.5 mm
  - Max Scan Size
    - 80 x 145 m ( $\emptyset$  x L)
- Capacity to scan rat vertebrae *in vivo* and large specimen (up to 80 mm in diameter)

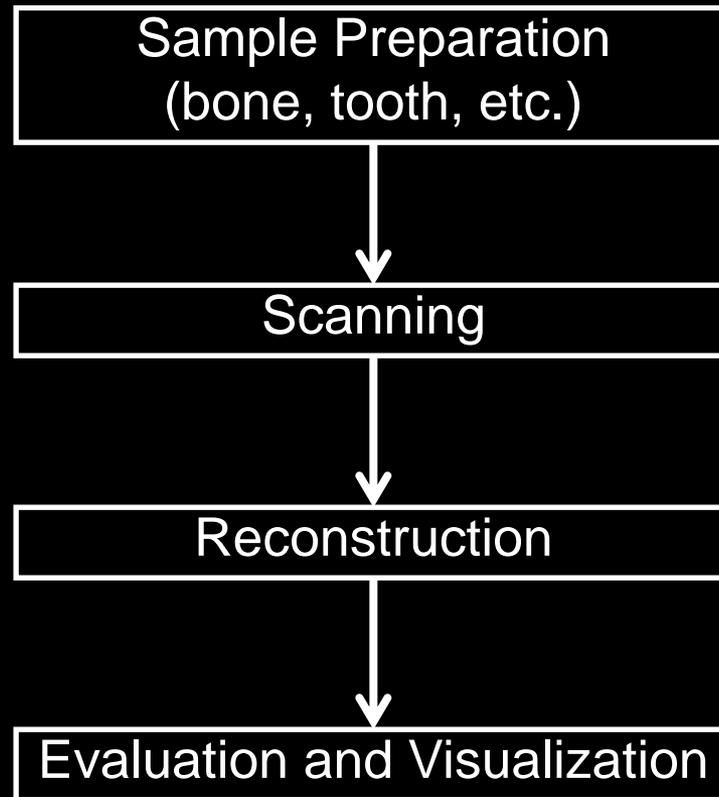


# Which Scanner Should I Use for My Study?

- vivaCT 40
  - Rodent study (rats and mice) study requiring in vivo scans (IACUC approval required)
  - When  $\mu$ CT 35 is fully occupied
- $\mu$ CT 35
  - Small specimen scans (under diameter of 35  $\mu$ m)
  - Mouse bone microstructure phenotyping **must** use  $\mu$ CT 35
- vivaCT 80
  - Rodent study (rats and mice) study requiring in vivo scans (IACUC approval required)
    - In vivo study of rat vertebrae and skull must use vivaCT 80 instead of vivaCT 40
  - Large specimen scans (above diameter of 40  $\mu$ m)
- vivaCT 75 and  $\mu$ CT 50 (VA location)
  - Only accessible to investigator with VA affiliations
  - $\mu$ CT 50: studies requiring high resolution characterization
  - vivaCT 75: large specimen scans (above diameter of 50  $\mu$ m)
- Extreme CT II
  - Clinical studies (IRB approval required)
  - Large specimen scans (above diameter of 80  $\mu$ m)
- **For consistent results, please use the same model of scanner for all samples/animals from the same study**



# $\mu$ CT Imaging



# Sample Preparation

- Specimens

- Dry samples

- Embedded in MMA ✓
    - Scanned in air (short scan time) for some soft tissue imaging ✓

- Wet samples

- Store in DI water ✓, saline ✓, PBS ✓, neutral buffer formalin ✓, ethonal ✓, etc.
    - Depend on the subsequent experiments (mechanical testing? Histology?)

Field of View (mm)	Projection /180°	Resolution (μm)
7.2	1000	3.5
12.3	1000	6
20.5	1000	10
30.7	1000	15
37.9	1000	18.5

- Image resolution

- Purpose of the study
  - Dimension of the sample

- Sample holders

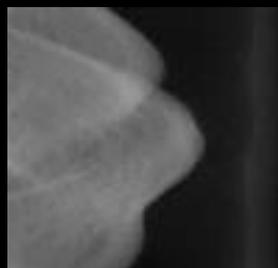
- 7-mm (3.5μm), 11.5-mm (6μm), 20-mm (10μm), 30-mm (15μm), 37-mm (18.5μm) holders
  - 15mL tube (10μm) or 50mL tube (15μm)

- **Position samples firmly in the holder (Very Important)**

- Use non-attenuating materials including foam, gauze, clay, paraffin, etc.

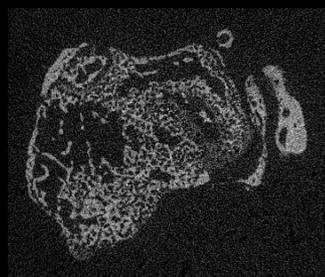


# From Scan to Results



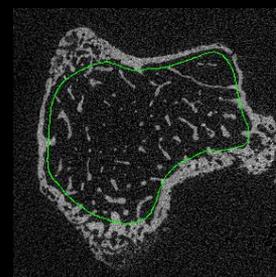
**.RSQ**  
RAW sequence data

Reconstruction →



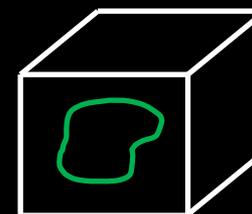
**.ISQ**  
Image sequence data

Contouring →



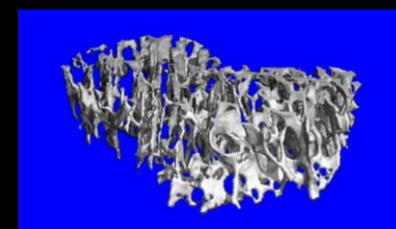
**.GOBJ**  
Graphical object

3D evaluation ↓



**.AIM**  
White box  
(gray scale)

Thresholding  
(Gauss) ↓



**\_SEG.AIM**  
Segmented object  
(binary file, black/white)

Calculating Morphometry ←



Evaluation Sheet

Trabecular Analysis

- \_TH.AIM
- \_SP.AIM
- \_TH.TXT
- \_SP.TXT

Cortical Analysis  
Midshaft Analysis

\_MOI.TXT

Final Results

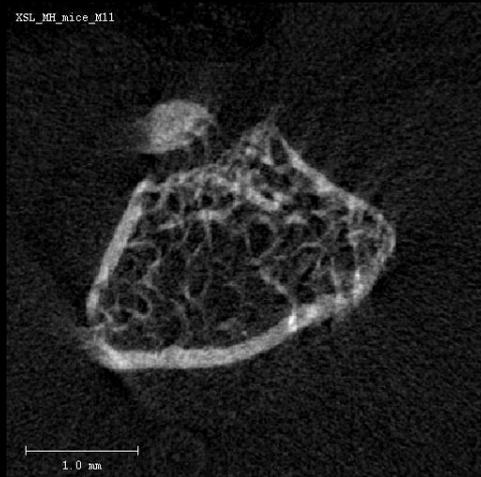
# After Scanning

- **Reconstruction**
  - Automatic reconstruction in Scanco (our) systems
- **Inspection of Images: Common Artifacts**
  - Motion artifact
  - Metal artifact
  - Ring artifact

# Motion Artifact

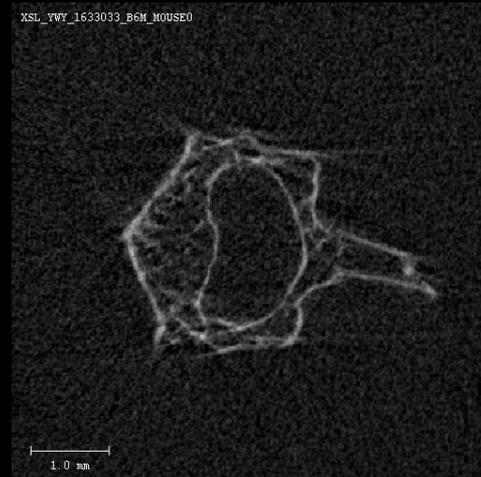
## Distal Femur

XSL\_MH\_mice\_M11

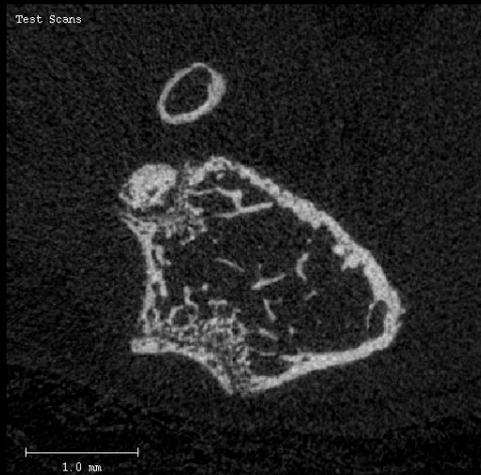


## L2 Vertebrae

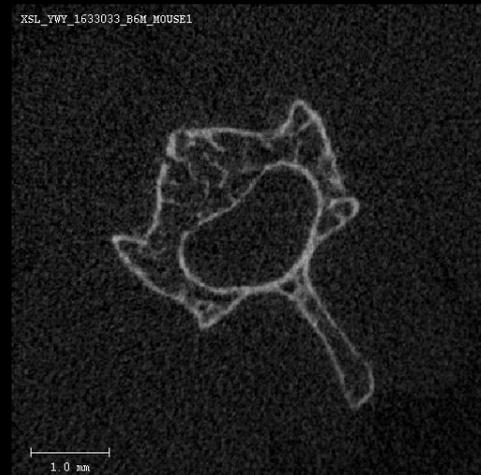
XSL\_FWY\_1633033\_B6M\_MOUSE0



Test Scans



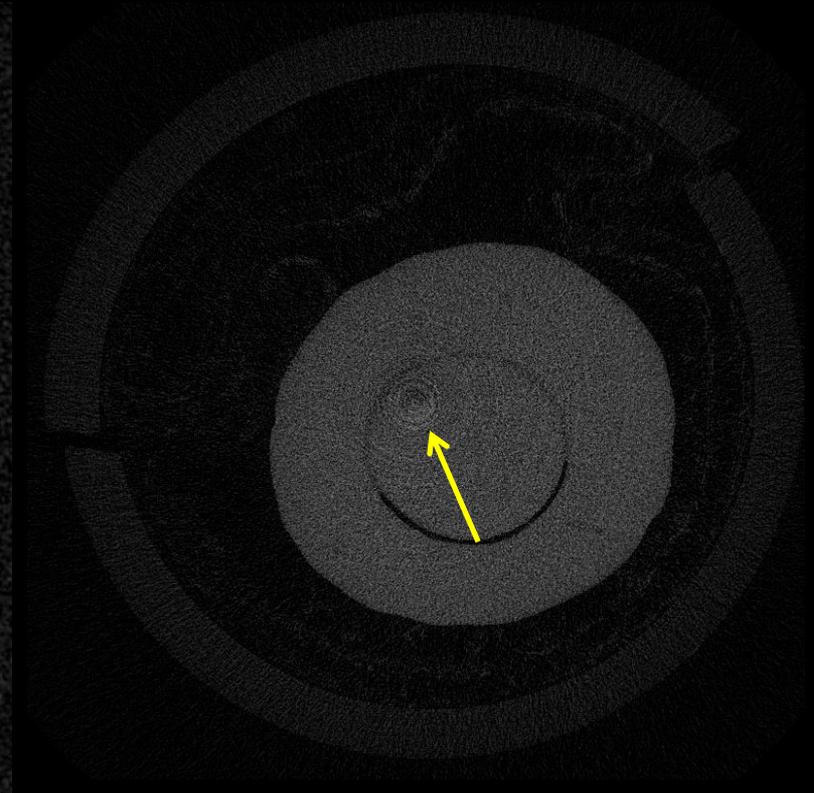
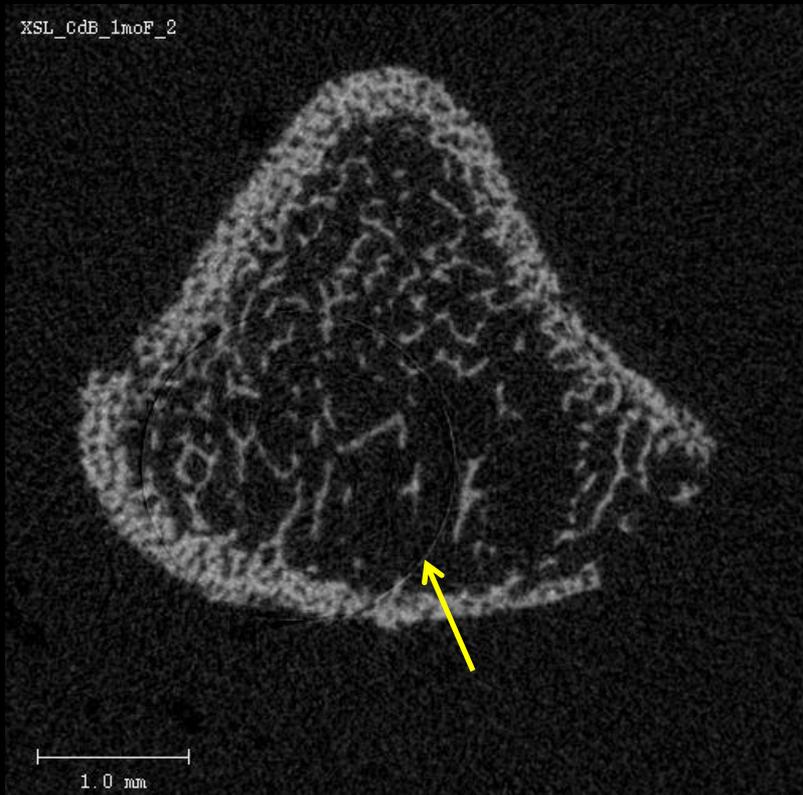
XSL\_FWY\_1633033\_B6M\_MOUSE1



# Metal Artifact



# Ring Artifact



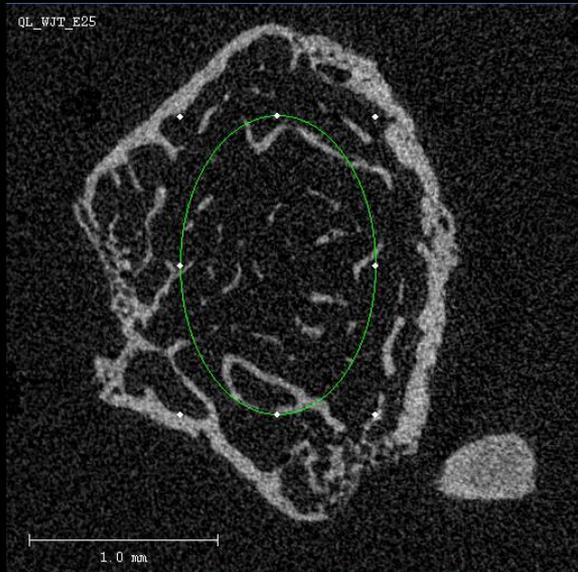
# After Scanning

- Reconstruction
  - Automatic reconstruction in Scanco (our) system
- Inspection of Images: Common Artifacts
  - Motion artifact
  - Metal artifact
  - Ring artifact
- **Image Processing**
  - Contouring
  - Filtration
  - Segmentation

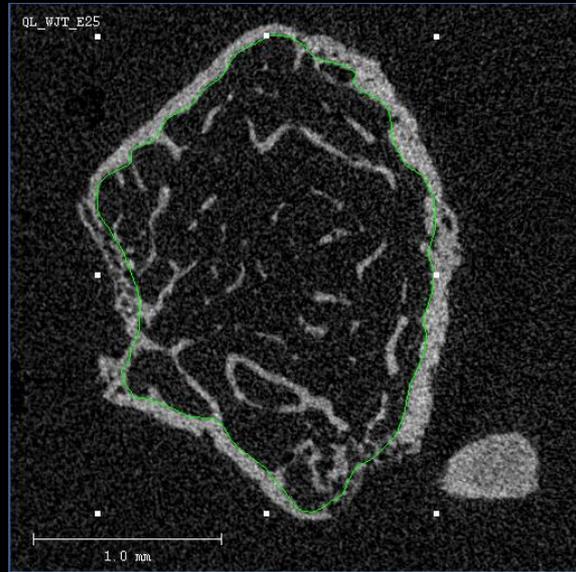
# Contouring

- Choose Volume of Interest (VOI)

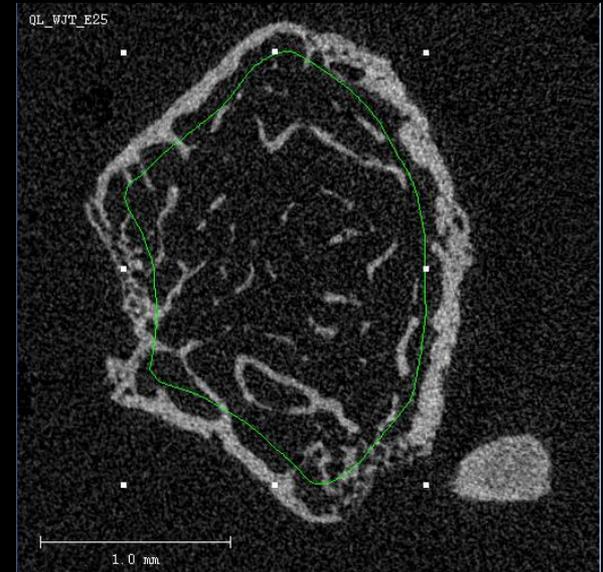
X



X

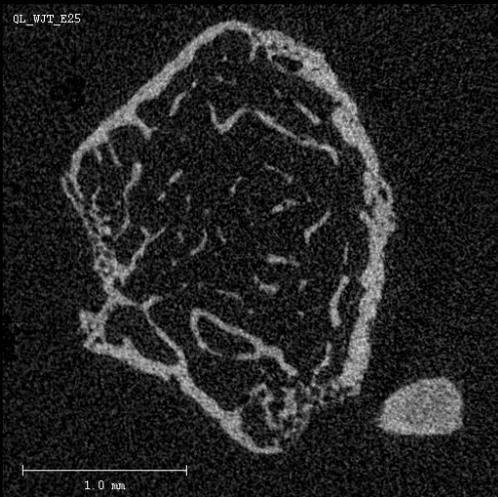


✓

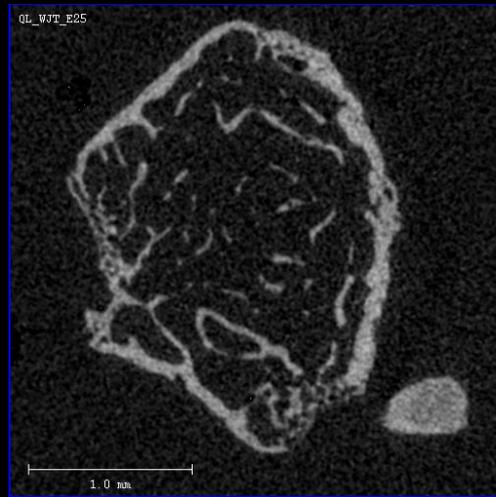


# Filtration

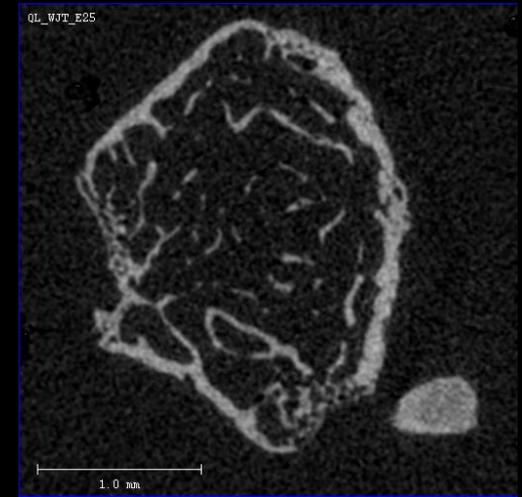
- Gaussian filter to suppress image noise
  - $\sigma$  : Extent of the kernel or degree of smoothing
  - Support: the size of the discrete kernel (# of neighboring voxels that contributes to the weighted mean)



unfiltered



$\sigma=1.2$ , support 2



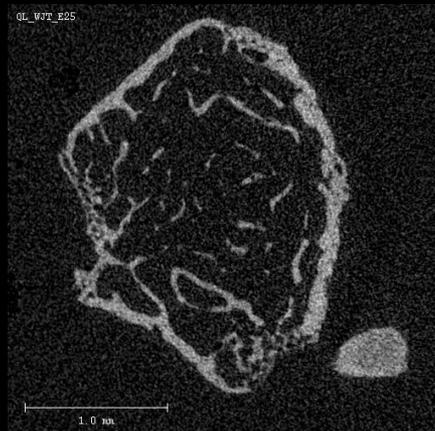
$\sigma=2$ , support 2

Recommended

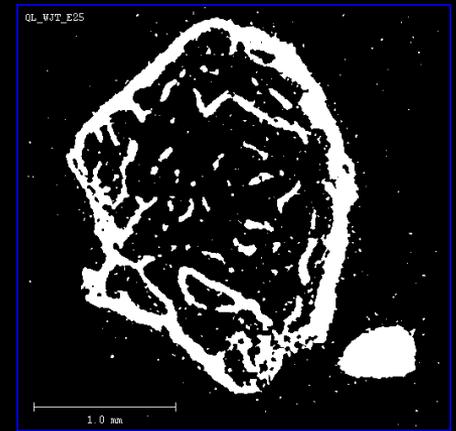
# Segmentation

- Separate the mineralized and non-mineralized tissue
  - **Low threshold:** a value ~200-400; Cut off the bone marrow and other non-mineralized tissue
  - **High threshold:** a fixed value of 1000; Cut off the white noise
  - **Recommended** low threshold: trabecular bone ~330; cortical bone ~360

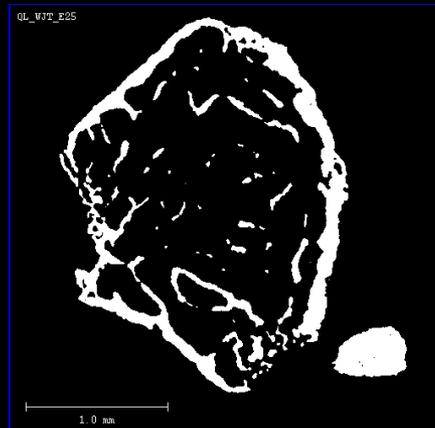
Grayscale



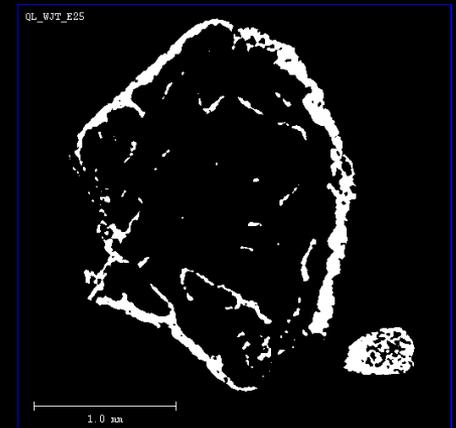
200-1000



Global Threshold  
330-1000



450-1000



Penn

# After Scanning

- Reconstruction
  - Automatic reconstruction in Scanco (our) system
- Inspection of Images: Common Artifacts
  - Motion artifact
  - Metal artifact
  - Ring artifact
- Image Processing
  - Contouring
  - Filtration
  - Segmentation
- **3D Evaluation**
  - Trabecular and Cortical bone density and microstructure



# Trabecular Bone Density and Microstructure

	Variable	Unit
TV	Total volume	mm <sup>3</sup>
BV	Bone volume	mm <sup>3</sup>
BV/TV	Bone volume fraction	%
SMI	Structure model index: 0 for parallel plates, 3 for cylindrical rods	
Conn.D	Connectivity Density	1/mm <sup>3</sup>
Tb.N	Trabecular number	1/mm
Tb.Th	Trabecular thickness	mm
Tb.Sp	Trabecular separation or spacing = marrow thickness	mm
BMD	Mean density values of everything within volume of interest	mg HA/cm <sup>3</sup>
TMD	Mean density of segmented region, thus only of what was considered bone	mg HA/cm <sup>3</sup>



# Trabecular Bone Density and Microstructure

- Scale

- Bone Volume Fraction (BV/TV)
- Trabecular Number (Tb.N)
- Trabecular Spacing (Tb.Sp)
- Trabecular Thickness (Tb.Th)  
(Hildebrand & Rueggsegger, 1995)

- Topology

- Connectivity (Conn.D) (Kinney and Labb, 1998; Kabel, *et al.* 1999)
- Structural Model Index (SMI) (Wehrl *et al.*, 2001; Hildebrand, *et al.* 1997)

- Orientation

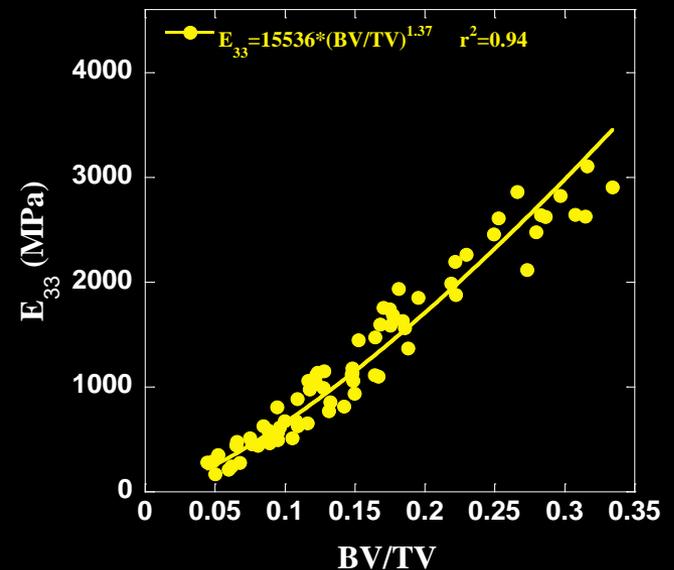
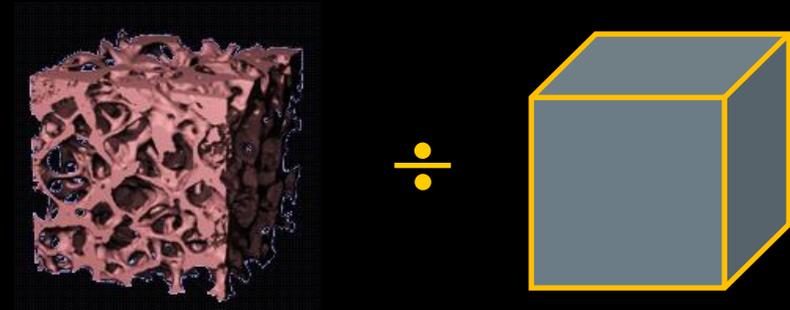
- Degree of Anisotropy (DA) (Cowin *et al.*, 1986)

- Bone Density

- Bone mineral density (BMD)
- Tissue mineral density (TMD)

Bone Volume Fraction (BV/TV)

$$\frac{\text{Bone Volume (BV)}}{\text{Total Volume (TV)}}$$



# Trabecular Bone Density and Microstructure

- Scale

- Bone Volume Fraction (BV/TV)
- Trabecular Number (Tb.N)
- Trabecular Spacing (Tb.Sp)
- Trabecular Thickness (Tb.Th)  
(Hildebrand & Ruegsegger, 1995)

- Topology

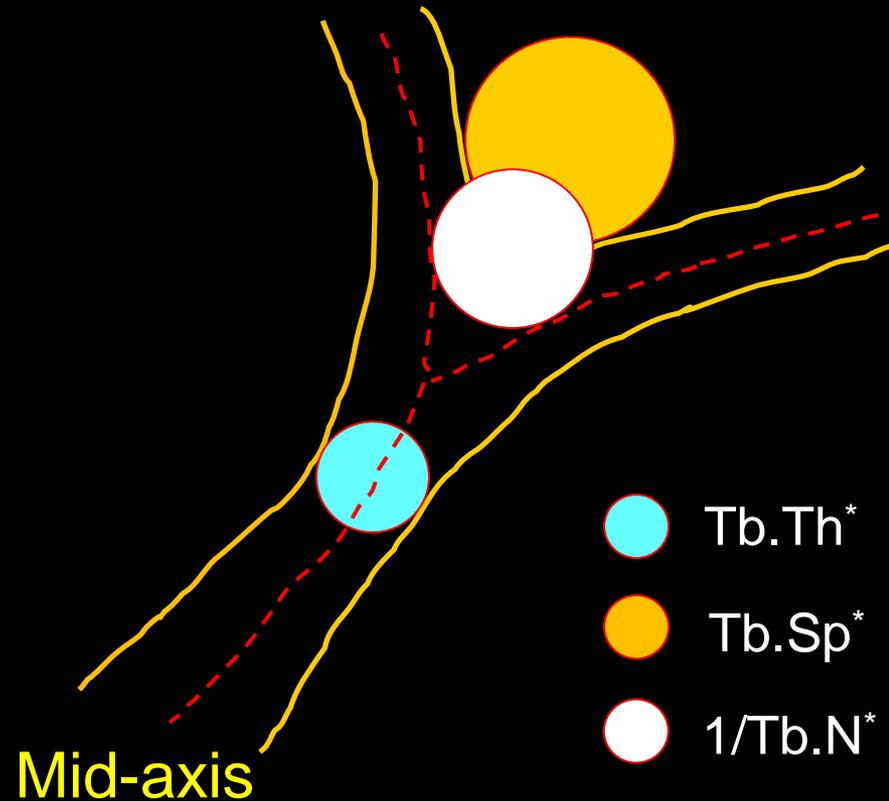
- Connectivity (Conn.D) (Kinney and Labb, 1998; Kabel, *et al.* 1999)
- Structural Model Index (SMI) (Wehrli *et al.*, 2001; Hildebrand, *et al.* 1997)

- Orientation

- Degree of Anisotropy (DA) (Cowin *et al.*, 1986)

- Bone Density

- Bone mineral density (BMD)
- Tissue mineral density (TMD)



# Trabecular Bone Density and Microstructure

- Topology

- Connectivity density (Conn.D) (Kinney and Labb, 1998; Kabel, *et al.* 1999)

- Indication of the integrity of trabecular bone network
    - Highly variable measurement outcome as it is sensitive to scan conditions, thresholding, etc
    - Limitation: plate perforation causes an increase in Conn.D

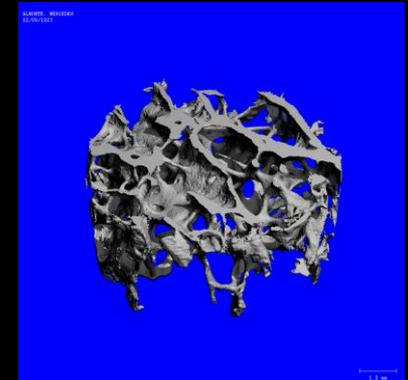
- Structural Model Index (SMI) (Wehrli *et al.*, 2001; Hildebrand, *et al.* 1997)

- Indication of plate-likeness of trabecular bone network
    - Increase of SMI indicates a conversion of trabecular bone from plate-like to rod-like, which is associated with decreased bone strength and increased fracture risk
    - SMI can be negative in trabecular bone evaluation, indicating a highly plate-like structure

- Orientation

- Degree of Anisotropy (DA) (Cowin *et al.*, 1986)

- Indication of how well trabeculae are aligned along the primary orientation within the network



A mix of plate- and rod-like structure

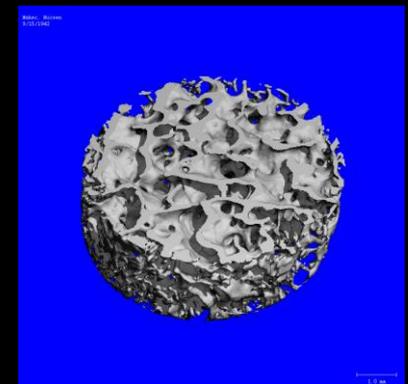


Plate-like structure



# Trabecular Bone Density and Microstructure

- Bone Density

- Bone mineral density (BMD)

- Total bone mineral content divided by the total volume (BMC/TV)
    - Mean density values of everything within the total volume
    - Rat tibia: ~300-400 mgHA/cm<sup>3</sup>
    - B6 WT Mice tibia: ~100-200 mgHA/cm<sup>3</sup>
    - B6 WT Mice femur: ~150-250 mgHA/cm<sup>3</sup>

- Tissue mineral density (TMD)

- Total bone mineral content divided by the bone volume (BMC/BV)
    - Mean density values of everything within the bone volume
    - Cortical bone: ~1000-1200 mgHA/cm<sup>3</sup>
    - Trabecular bone: ~800-1000 mgHA/cm<sup>3</sup>



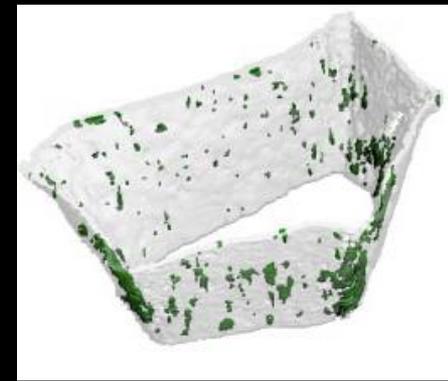
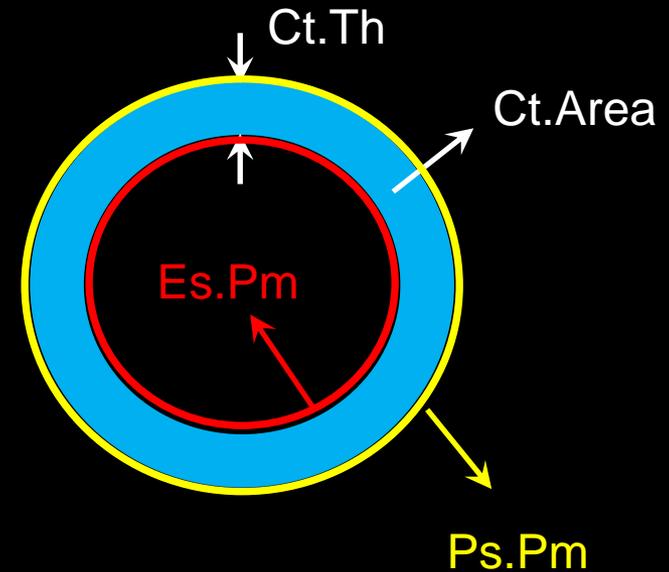
# Cortical Bone Morphometry

	Variable	Unit
Ct.Area	Cortical Area	mm <sup>2</sup>
Ct.Th	Cortical thickness	mm
Ct.Po	Cortical Porosity	%
pMOI	Polar moment of inertia	mm <sup>4</sup>
Ps.Pm	Periosteal perimeter	mm
Es.Pm	Endocortical perimeter	mm
BMD	Mean density values of everything within volume of interest	mg HA/cm <sup>3</sup>
TMD	Mean density of segmented region, thus only of what was considered bone	mg HA/cm <sup>3</sup>



# Cortical Bone Morphometry

- Cortical thickness (Ct.Th)
  - Thickness of the cortex
- Cortical area (Ct.Area)
  - Mean surface area of the cortex
  - Bone volume divided by the bone length
- Periosteal perimeter (Ps.Pm)
- Endosteal perimeter (Es.Pm)
- Cortical porosity (Ct.Po)
  - Percent of void volume over bone volume
  - ~1-10%



Burghardt *et al.* J Clin Endocrinol Metab 2010

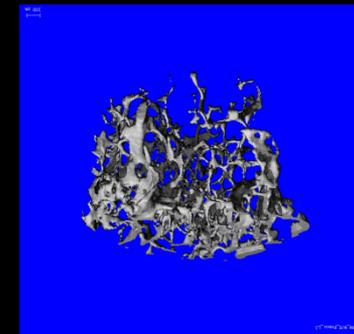
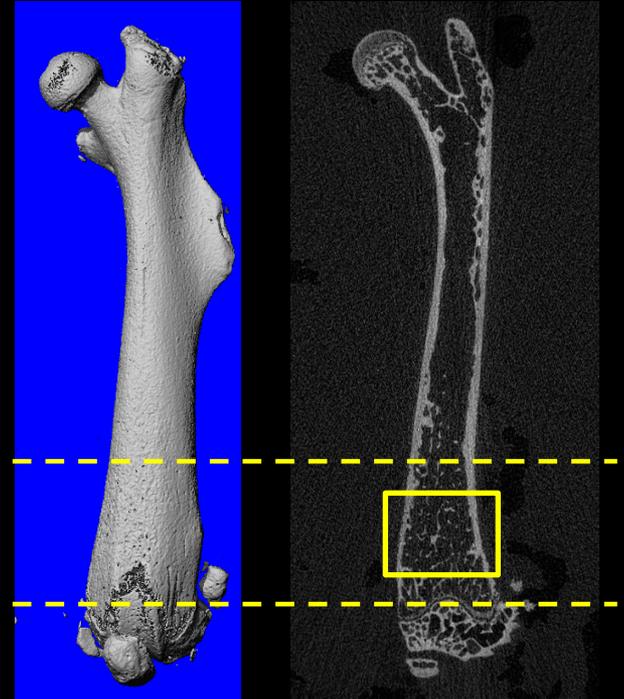
# Post-Processing

- Visualization – 3D Rendering
- Export your scans
  - DICOM format
  - TIFF format
- Software (free) for post-processing
  - Scanco Image Process Language
  - ImageJ – BoneJ (Analysis)
  - MicroView (Analysis)
  - CTAn (Bruker) (Analysis)
  - OsiriX (Segmentation, Visualization, Analysis)
  - ITK-SNAP (Segmentation, Visualization)
  - Slicer (Segmentation, Registration)
- Export for 3D printing
  - STL format



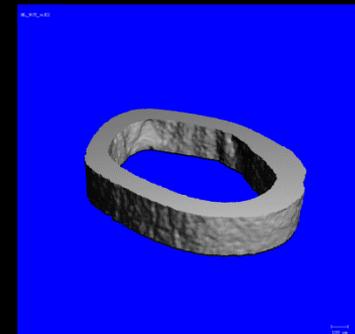
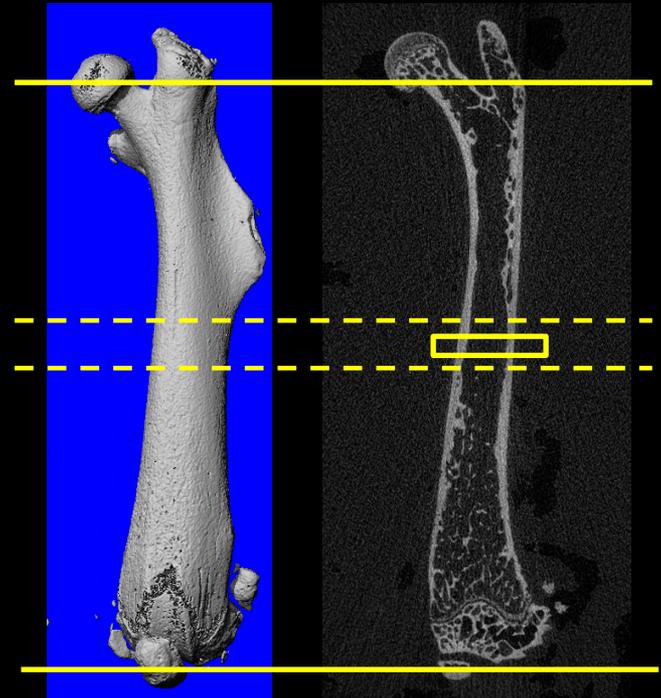
# Phenotyping

- Mouse trabecular bone (long bone)
  - Distal Femur, proximal tibia
  - 6  $\mu\text{m}$  isotropic voxel size
  - Scan Region
    - 1-2 stacks, ~210-420 slices
  - Average scan time: 24-48 mins
  - Analysis region
    - 100-200 slices, 0.5-1mm from the growth plate
  - Outcome measures: BV/TV, Tb.Th, Tb.N, Tb.Sp, SMI, Conn.D, BMD, TMD



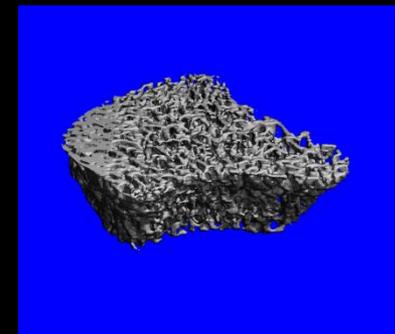
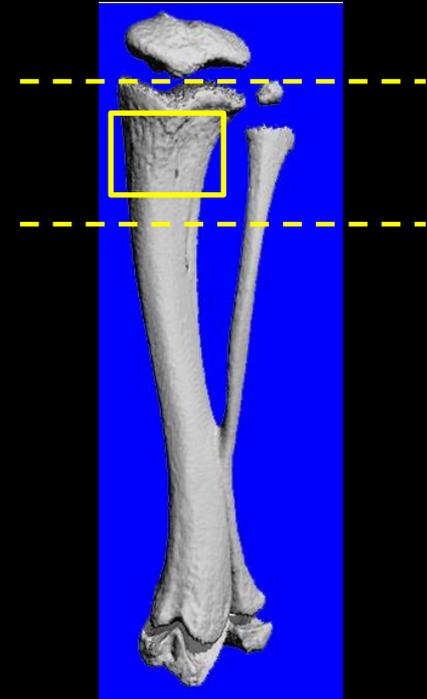
# Phenotyping

- Mouse cortical bone (long bone)
  - Midshaft of tibia or femur
  - 6-10  $\mu\text{m}$  isotropic voxel size
  - Scan Region
    - 1 stacks, ~210 slices
  - Average scan time: 24 mins
  - Analysis region
    - Middle 50 slices
  - Outcome measures: Ct.Area, Ct.Th, pMOI, Ct.Po, TMD



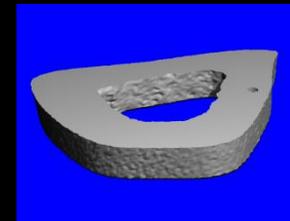
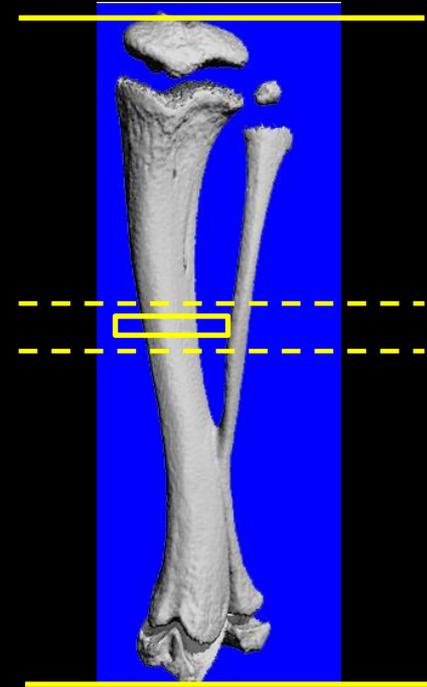
# Phenotyping

- Rat trabecular bone (long bone)
  - Proximal tibia
  - 6-10  $\mu\text{m}$  isotropic voxel size
  - Scan Region
    - 2 stacks,  $\sim 420$  slices
  - Average scan time: 24-48 mins
  - Analysis region
    - $\sim 200$  slices, 1-2mm distal to the growth plate
  - Outcome measures: BV/TV, Tb.Th, Tb.N, Tb.Sp, SMI, Conn.D, TMD



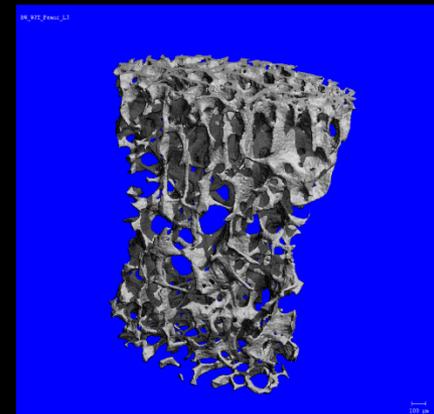
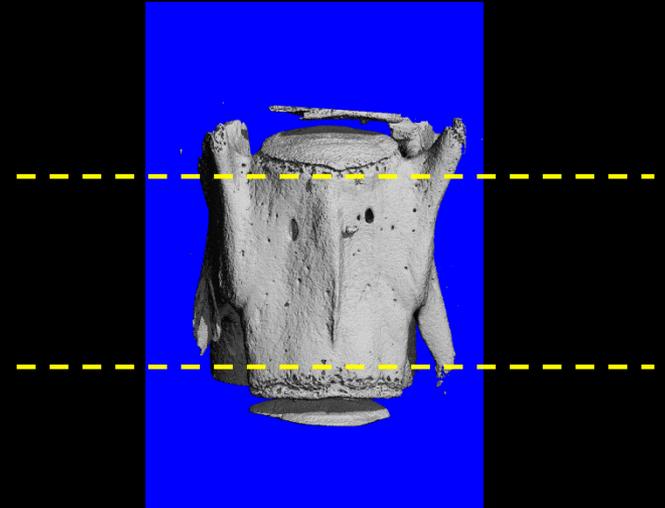
# Phenotyping

- Rat cortical bone (long bone)
  - Midshaft of tibia or femur
  - 6-10  $\mu\text{m}$  isotropic voxel size
  - Scan Region
    - 1 stacks, ~210 slices
  - Average scan time: 24 mins
  - Analysis region
    - Middle 50 slices
  - Outcome measures: Ct.Area, Ct.Th, pMOI, Ct.Po, TMD



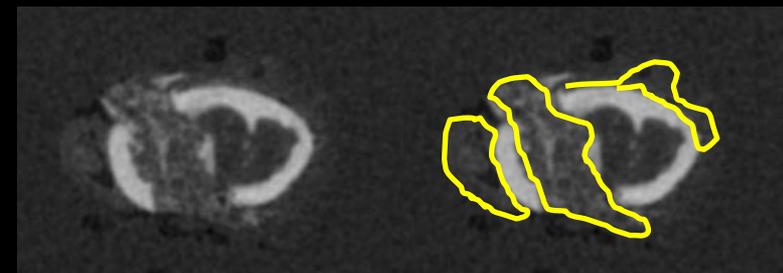
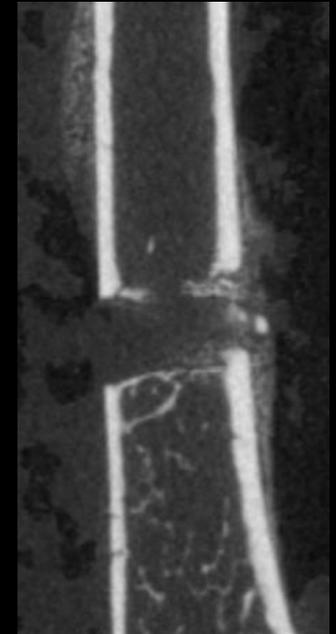
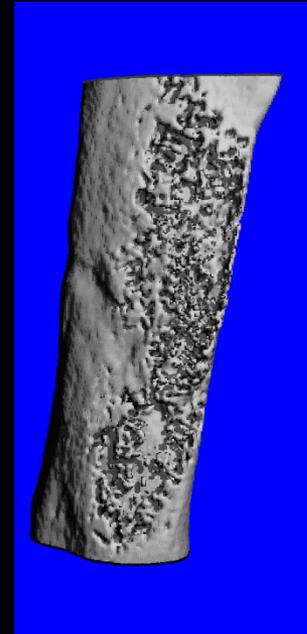
# Phenotyping

- Mouse/Rat vertebral trabecular bone
  - Lumbar vertebra L1-L4
  - 6  $\mu\text{m}$  (mouse) / 6-10  $\mu\text{m}$  (rat) isotropic voxel size
  - Scan Region
    - Between end plates
  - Average scan time: 48-72 mins
  - Analysis region
    - Between two end plates
    - Middle 150-200 slices
  - Outcome measures: BV/TV, Tb.Th, Tb.N, Tb.Sp, SMI, Conn.D, BMD, TMD



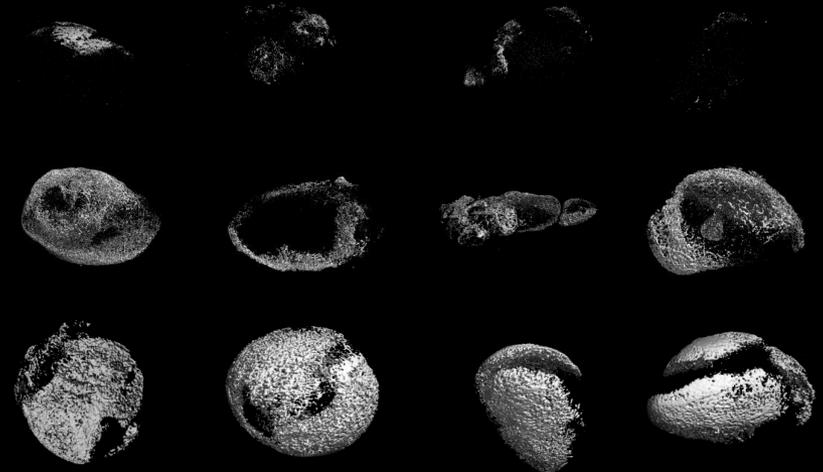
# Fracture Healing

- 6-10  $\mu\text{m}$  isotropic voxel size
- Scan region
  - Whole bone
- Average scan time
  - Scan vertically: 1-2 hours
  - Scan horizontally: 15-30 mins
- Analysis region
  - whole healing region excluding cortical and trabecular bone
- Outcome measures
  - BV: Callus Volume
  - Avg callus area =  $\text{Callus Volume} / \text{Total Length}$



# Scaffolds

- PLA, PGA, PCL, etc.
  - 6 -10  $\mu\text{m}$  isotropic voxel size
  - Scan Region:
    - Whole scaffold
  - Average scan time:
    - 0.5-2 hours
  - Analysis Region:
    - Whole scaffold
  - Outcome measure:
    - Total mineralized tissue content ( $\text{BV} * \text{TMD}$ )
    - Total mineralized tissue volume (BV)



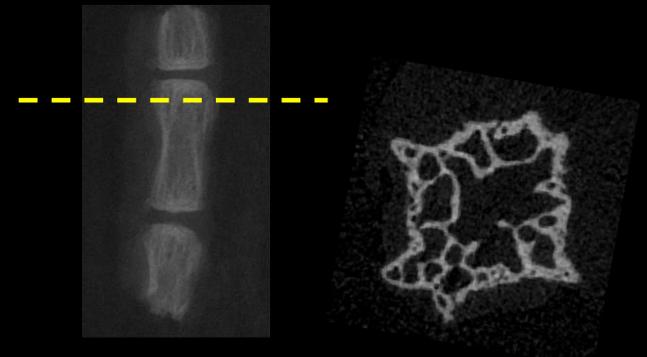
Courtesy of Dr. Masahiro and Dr. Kenta Uchibe



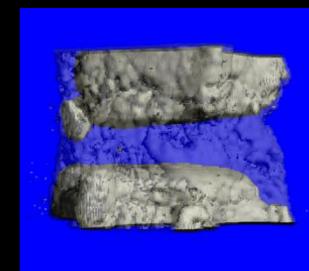
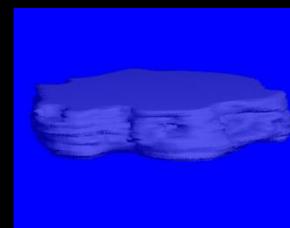
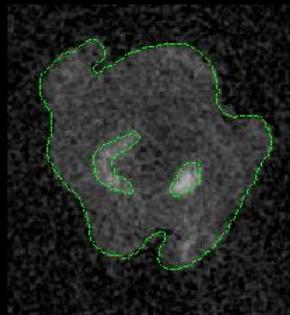
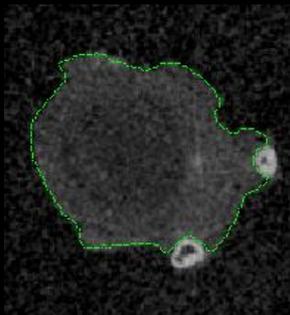
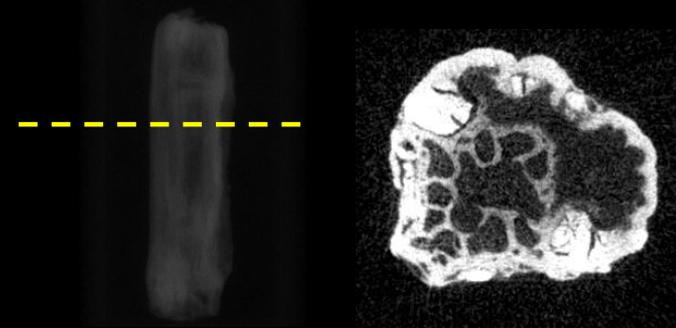
# Soft Tissue $\mu$ CT Imaging

- Contrast-Enhanced  $\mu$ CT Imaging
  - Mouse Intervertebral disc
  - 18.5 $\mu$ m or lower voxel size
  - Spine segment scanned before and after Lugols staining
  - Scan Region:
    - Whole segment
  - Average scan time:
    - 0.5-2 hours
  - Outcome measures
    - Disc thickness (Tb.Th)

Before

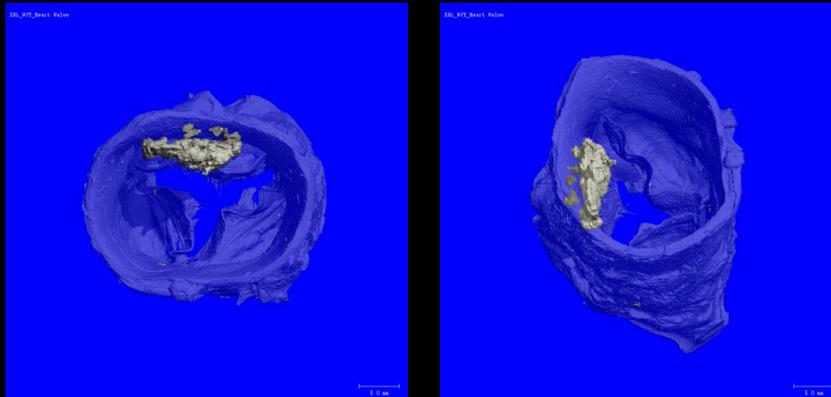


After



# Soft Tissue $\mu$ CT Imaging

- Soft tissue scanned in air
  - Heart Valve
  - Larynx Branch



Heart Valve



Larynx Branch

# Questions?