

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

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

What is μ 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 (μ FE) models to estimate the mechanical properties of bone van Rietbergen+1998, Schulte+2011



Available Scanners at PCMD μ CT Imaging Core

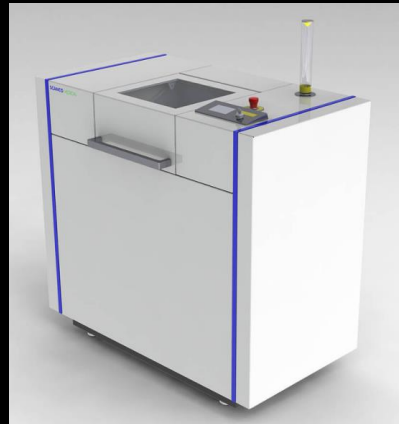
- Specimen μ CT

- μ CT 35
- μ CT 50
- μ CT 45 *new!*



- *In Vivo* μ CT

- vivaCT 40
- vivaCT 75
- vivaCT 80 *new!*



- Clinical μ CT

- XtremeCT II



Penn

PCMD *ex vivo* μ CT Scanners

	μ CT 35	μ CT 45	μ CT 50
Use	Specimen	Specimen	Specimen
X-Ray Source	30 - 70 kVp	20 - 90 kVp	30 - 90 kVp
Max Scan Size	37.9 x 120 mm ($\varnothing$ x L)	50 x 120 mm ($\varnothing$ x L)	50 x 120 mm ($\varnothing$ x L)
Max Specimen Size	75.8 x 140 mm ($\varnothing$ x L)	90 x 120 mm ($\varnothing$ x L)	100 x 160 mm ($\varnothing$ x L)
Best image voxel size	3.5 μ m ($\varnothing$:7 mm)	2.5 μ m ($\varnothing$: 5 mm)	1.5 μ m ($\varnothing$: 3 mm)
Location	McKay Lab	McKay Lab	VA Hospital

PCMD *in vivo* μ CT Scanners

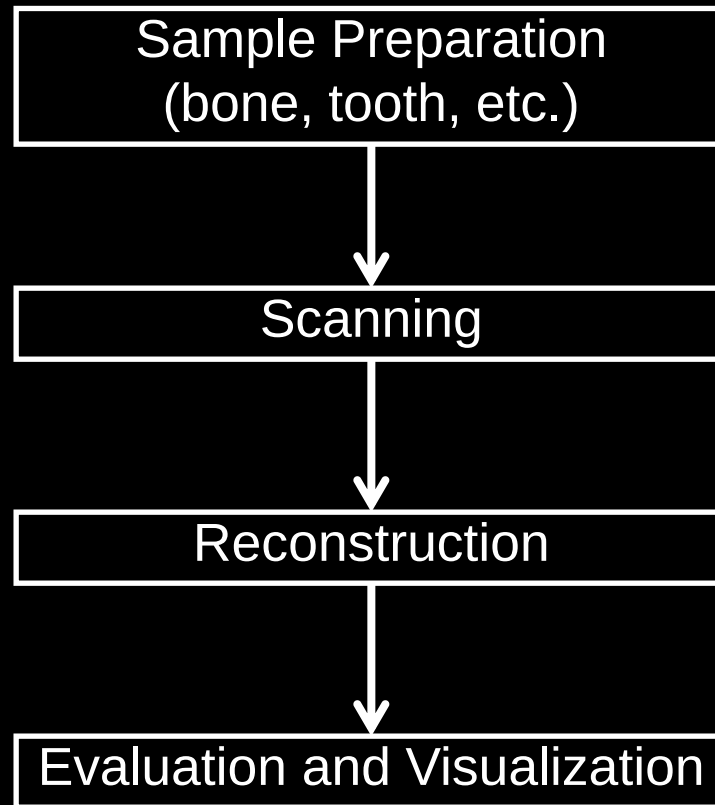
	vivaCT 40	vivaCT 80	vivaCT 75	XtremeCT II
Use	In Vivo Rodent	In Vivo Rodent	In Vivo Rodent	Clinical
X-Ray Source	30 - 70 kVp	30 - 70 kVp	30 - 70 kVp	68 kVp
Max Scan Size	38.9 x 145 mm ($\varnothing$ x L)	80 x 145 mm ($\varnothing$ x L)	79.9 x 145 mm ($\varnothing$ x L)	140 x 200 mm ($\varnothing$ x L)
Max Specimen Size	80 x 500 mm ($\varnothing$ x L)	90 x 500 mm ($\varnothing$ x L)	80 x 500 mm ($\varnothing$ x L)	170 mm ($\varnothing$)
Best image voxel size	10.5 μ m ($\varnothing$:21.5 mm)	10.4 μ m ($\varnothing$:21.5 mm)	19 μ m ($\varnothing$: 38.9 mm)	60 μ m
Location	McKay Lab	McKay Lab	VA Hospital	CHOP CTRC nutrition and growth lab

Which Scanner Should I Use for My Study?

- μ CT 35 and μ CT 45
 - Small specimen scans (under diameter of 35 μ m)
 - Mouse bone microstructure phenotyping **must** use μ CT 35/45
- vivaCT 40
 - Rodent study (rats and mice) requiring in vivo scans (IACUC approval required)
 - When μ CT 35/45 is fully occupied
- vivaCT 80
 - Rodent study (rats and mice) 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 50 μ m)
- vivaCT 75 and μ CT 50 (VA location)
 - Only accessible to investigators with VA affiliations
 - μ CT 50: studies requiring high resolution characterization
 - vivaCT 75: large specimen scans (above diameter of 50 μ m)
- Extreme CT II
 - Clinical studies (IRB approval required)
 - Large specimen scans (above diameter of 80 μ m)
- **For consistent results, please use the same model of scanner for all samples/animals from the same study**



μ CT Imaging



Sample Preparation

- Specimens
 - Dry samples
 - Embedded in MMA ✓
 - Scanned in air (short scan time) for special purposes of soft tissue imaging ✓
 - Wet samples
 - Stored in DI water ✓, saline ✓, PBS ✓, neutral buffer formalin ✓, ethonol ✓, etc.
 - Depend on the subsequent experiments (mechanical testing? Histology?)

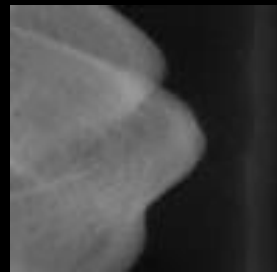
- Image resolution
 - Purpose of the study
 - Dimension of the sample

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

- Sample holders
 - 7-mm (3.5 μm), 11.5-mm (6 μm), 20-mm (10 μm), 30-mm (15 μm), 37-mm (18.5 μm) holders
 - 15mL tube (10 μm) or 50mL tube (15 μm)
- Position samples firmly in the holder (Very Important)
 - Use non-attenuating materials including foam, gauze, clay, paraffin, etc.

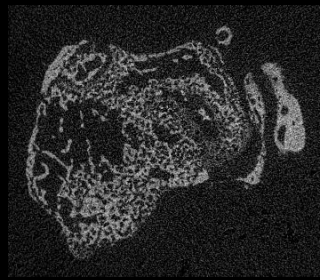


From Scan to Results



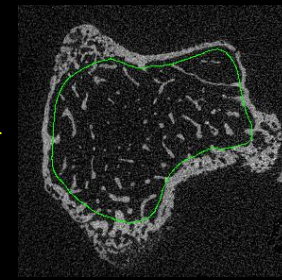
.RSQ
RAW sequence data

Reconstruction →



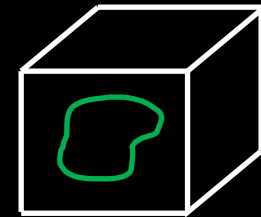
.ISQ
Image sequence data

Contouring →



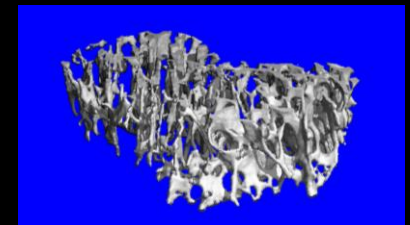
.GOBJ
Graphical object

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



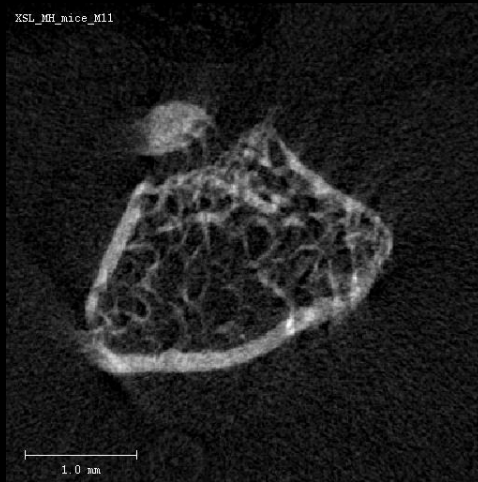
Penn

After Scanning

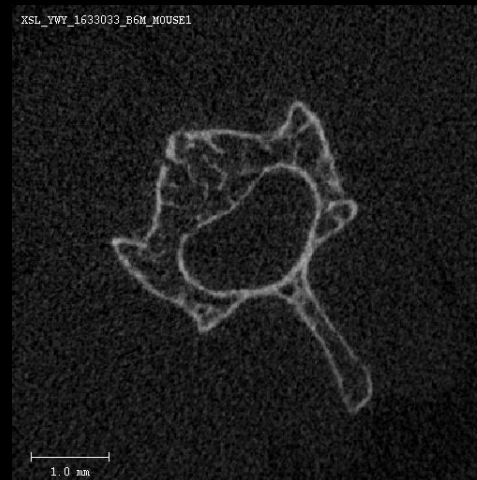
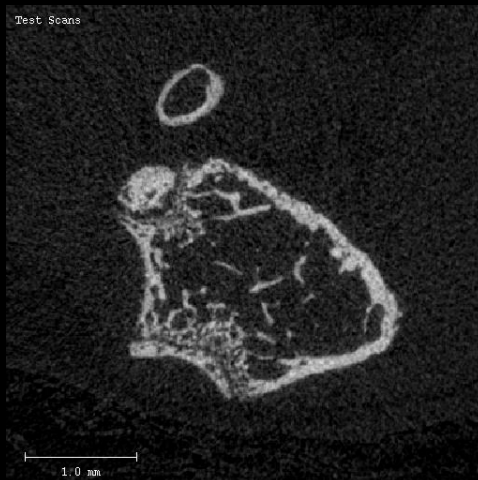
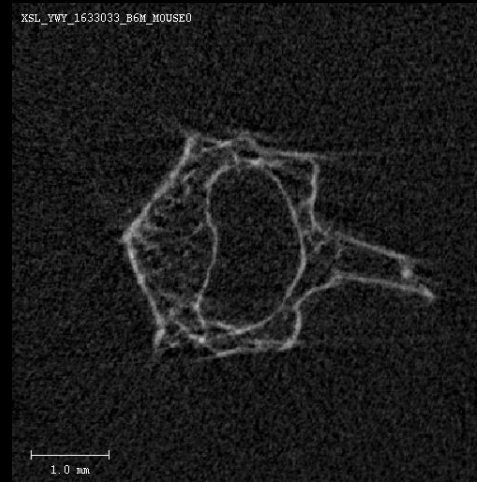
- **Reconstruction**
 - Automatic reconstruction in Scanco (our) systems
- **Inspection of Images: Common Artifacts**
 - Motion artifact
 - Metal artifact
 - Ring artifact

Motion Artifact

Distal Femur



L2 Vertebrae

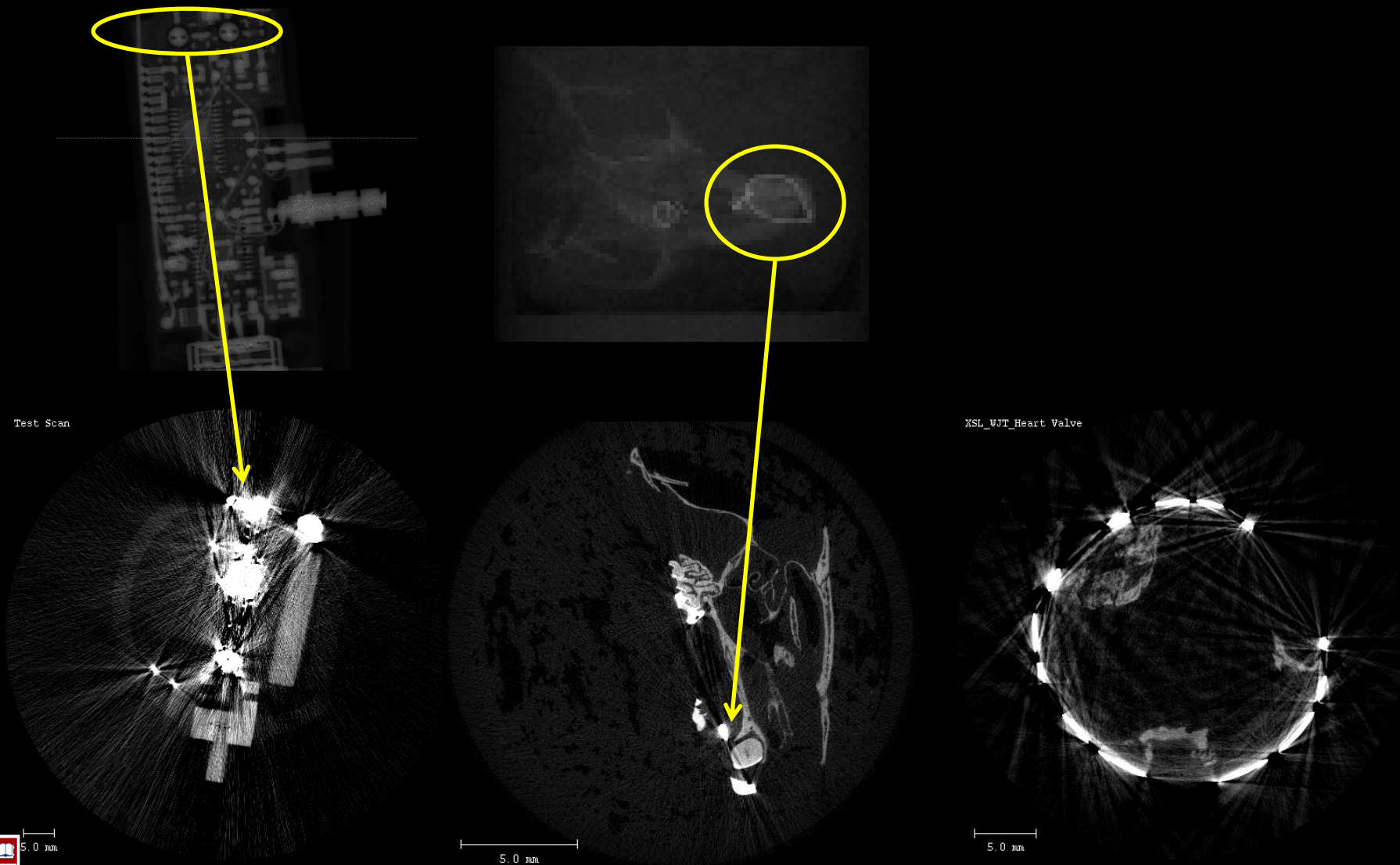


Sometimes re-scans required

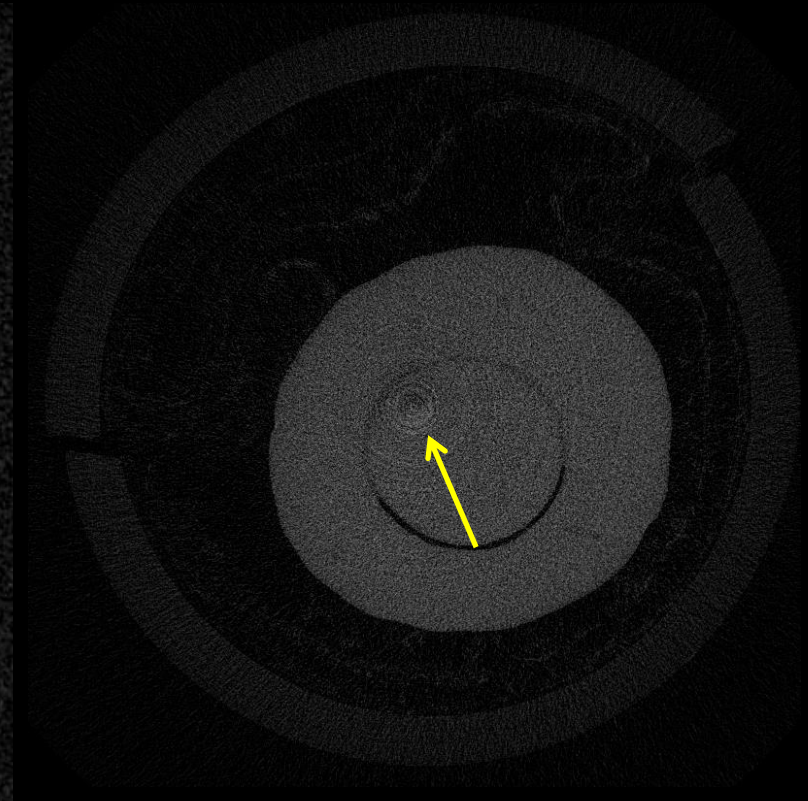
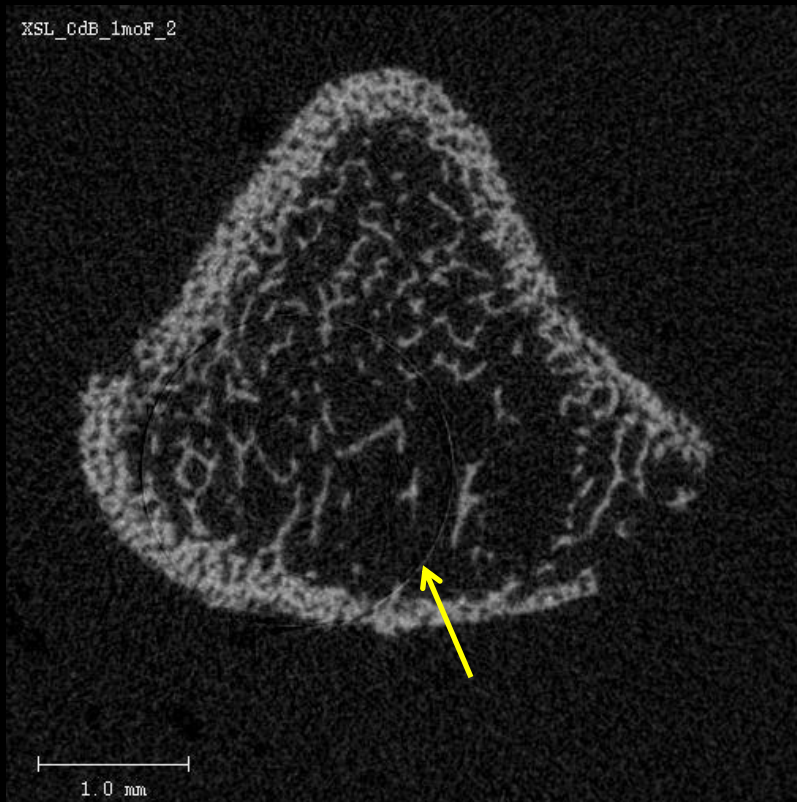


Penn

Metal Artifact



Ring Artifact



Report to the core manager with sample and measurement #



Penn

After Scanning

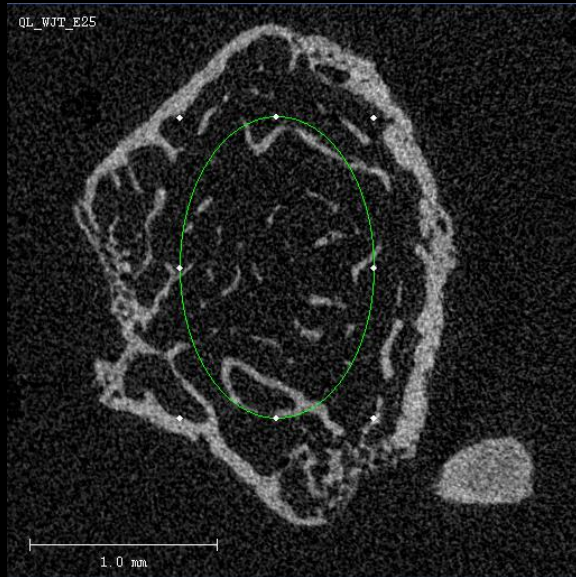
- Reconstruction
 - Automatic reconstruction in Scanco (our) system
- Inspection of Images: Common Artifacts
 - Motion artifact
 - Metal artifact
 - Ring artifact
- Image Processing
 - Contouring
 - Filtration
 - Segmentation



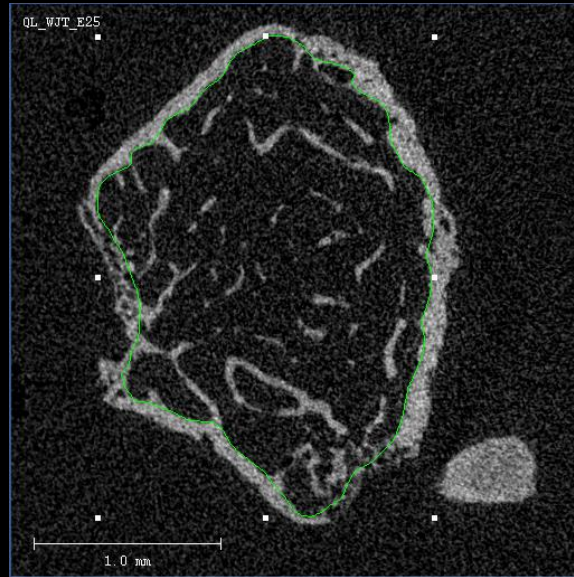
Contouring

- Choose Volume of Interest (VOI)

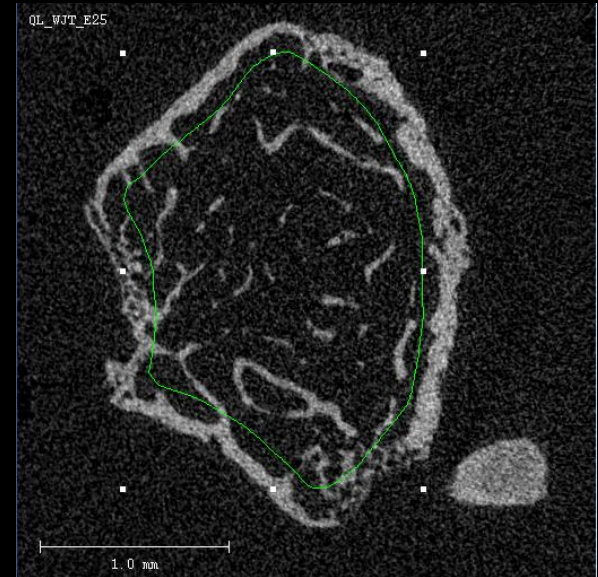
X



X

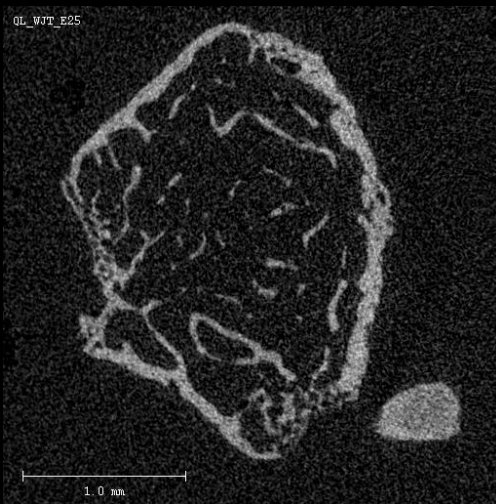


✓

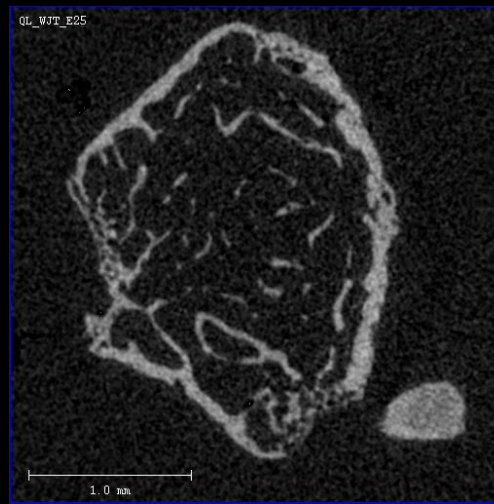


Filtration

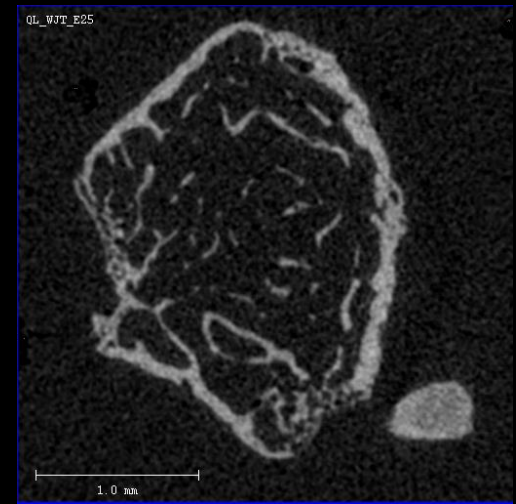
- Gaussian filter to suppress image noise
 - σ : Extent of the kernel or degree of smoothing
 - Support: the size of the discrete kernel (# of neighboring voxels that contributes to the weighted mean)



unfiltered



$\sigma=1.2$, support 2



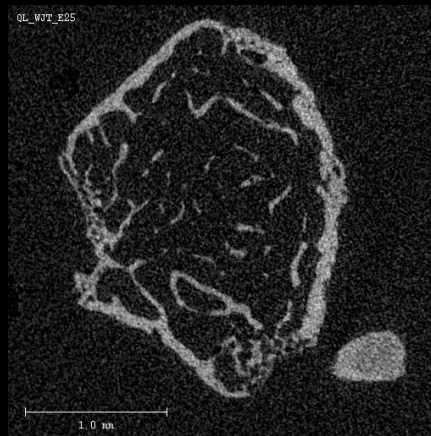
$\sigma=2$, support 2

Recommended

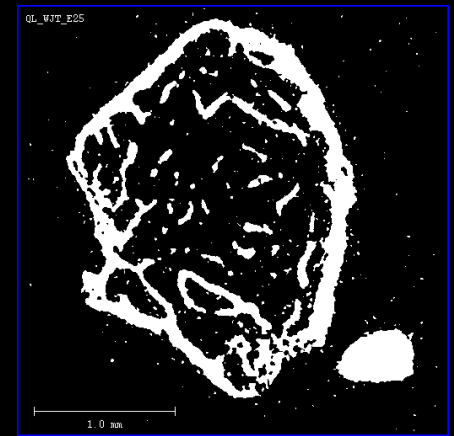
Segmentation

- 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
 - **Recommendation:** low threshold: trabecular bone ~330; cortical bone ~360

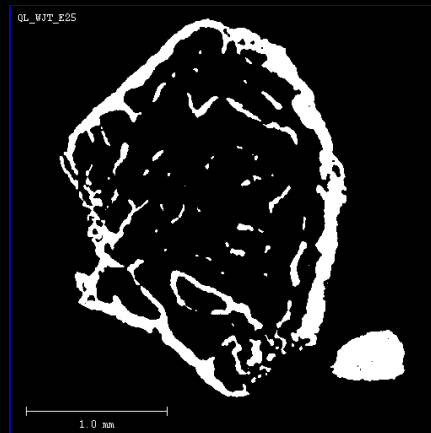
Grayscale



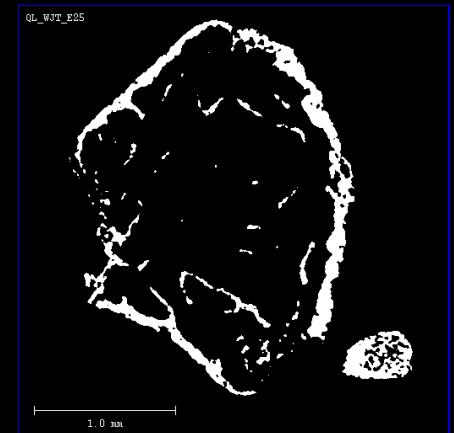
200-1000



Global Threshold
330-1000



450-1000



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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 ³
BV	Bone volume	mm ³
BV/TV	Bone volume fraction	%
SMI	Structure model index: 0 for parallel plates, 3 for cylindrical rods	
Conn.D	Connectivity Density	1/mm ³
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 ³
TMD	Mean density of segmented region, thus only of what was considered bone	mg HA/cm ³



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)

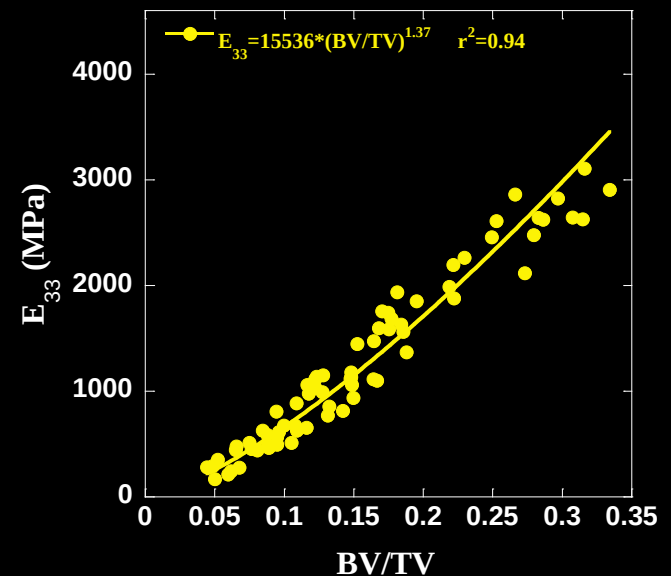
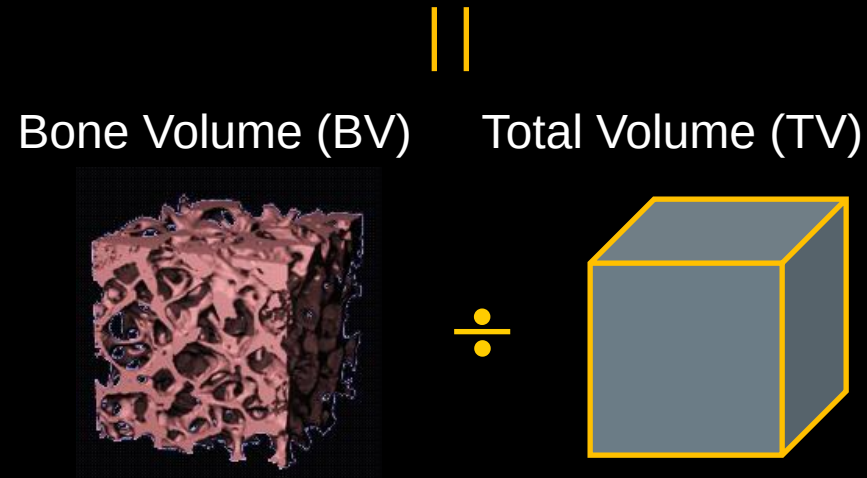
- Orientation

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

- Bone Density

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

Bone Volume Fraction (BV/TV)



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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

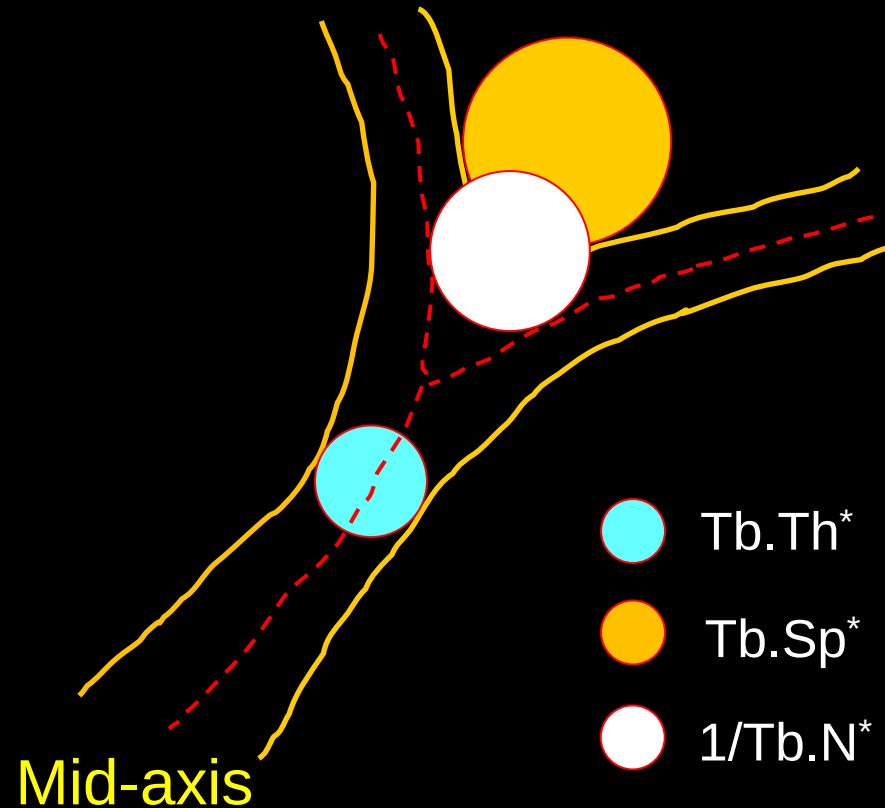
- Connectivity (Conn.D) (Kinney and Labb, 1998; Kabel, *et al.* 1999)
- Structural Model Index (SMI) (Wehrli *et al.*, 2001; Hildebrand, *et al.* 1997)

- Orientation

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

- Bone Density

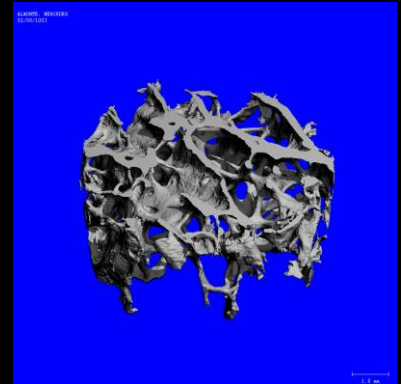
- Bone mineral density (BMD)
- Tissue mineral density (TMD)



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
 - A 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 the 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



A mix of plate- and rod-like structure

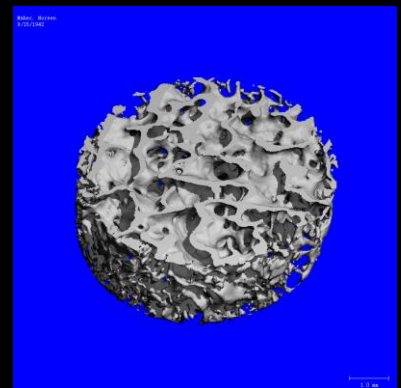


Plate-like structure

- Orientation

- Degree of Anisotropy (DA) (Cowin *et al.*, 1986)
 - Indication of how well trabeculae are aligned along the primary orientation within the network



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/cm³
 - B6 WT Mice tibia: ~100-200 mgHA/cm³
 - B6 WT Mice femur: ~150-250 mgHA/cm³
 - 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/cm³
 - Trabecular bone: ~800-1000 mgHA/cm³

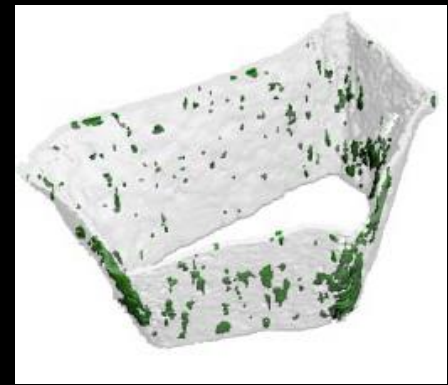
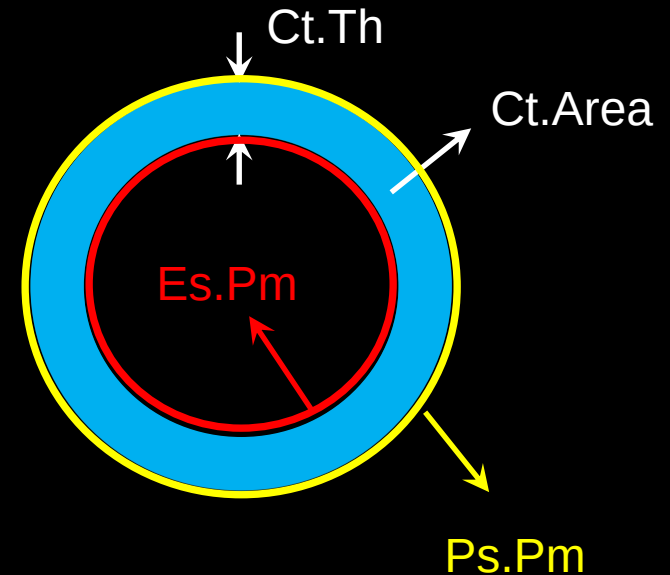
Cortical Bone Morphometry

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



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

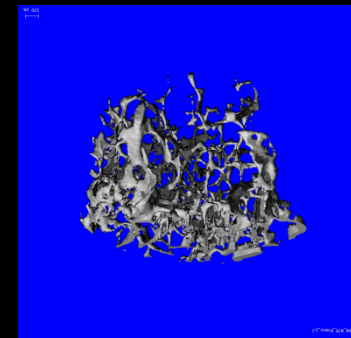
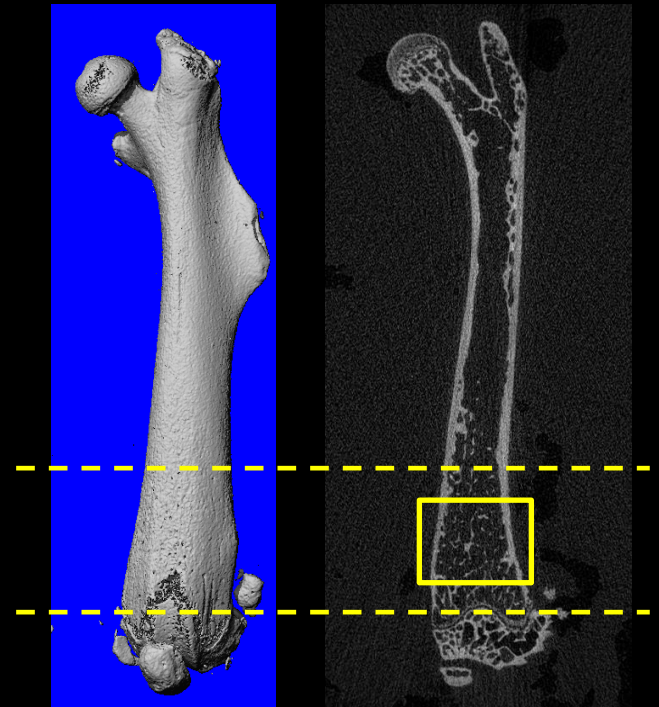
Post-Processing

- 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



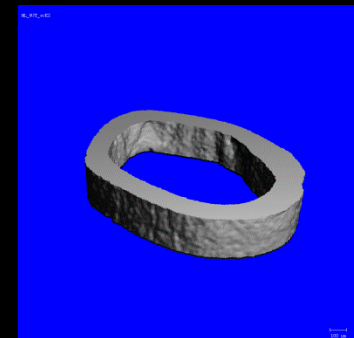
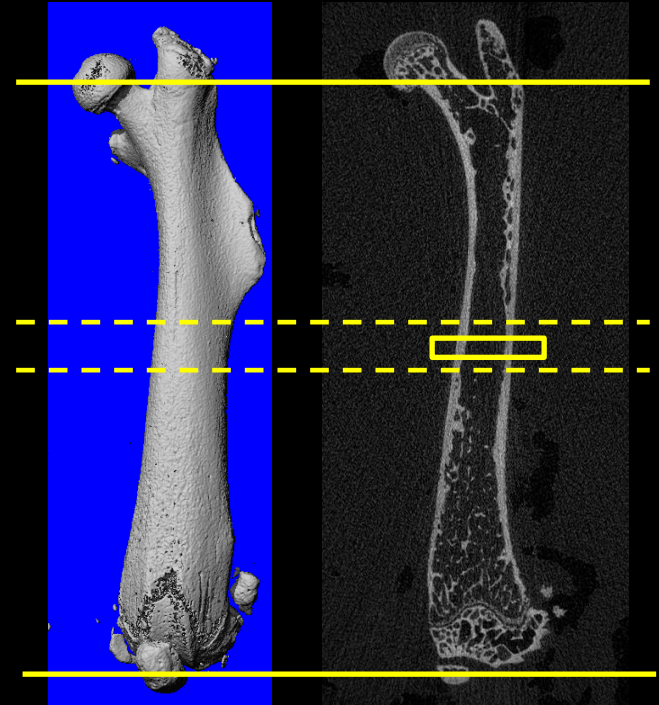
Phenotyping

- Mouse trabecular bone (long bone)
 - Distal femur, proximal tibia
 - 6 μm 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



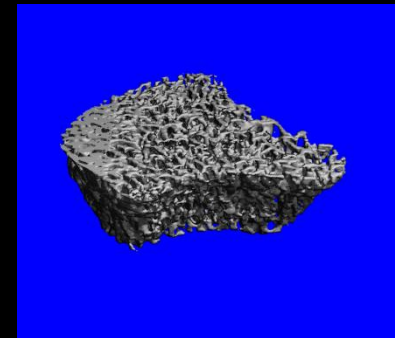
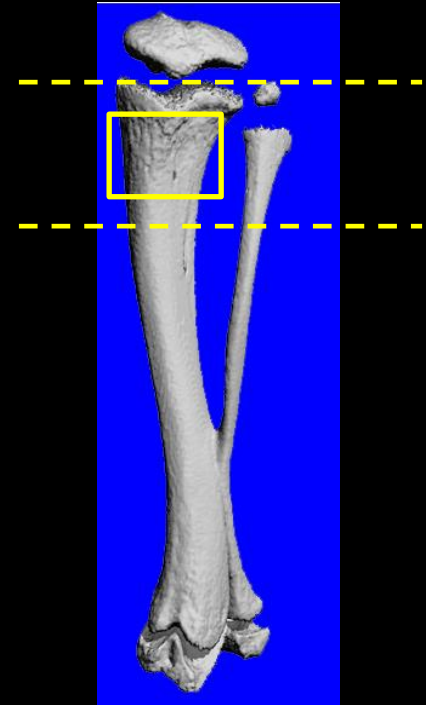
Phenotyping

- Mouse cortical bone (long bone)
 - Midshaft of tibia or femur
 - 6-10 μm 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



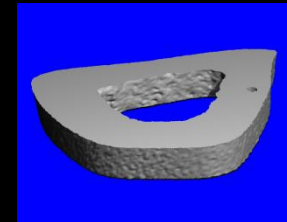
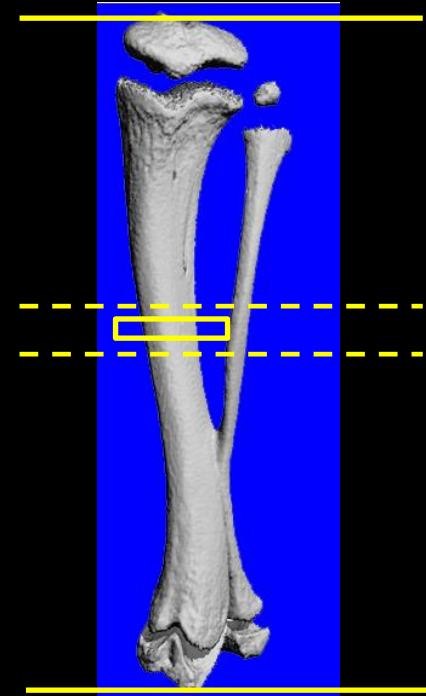
Phenotyping

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



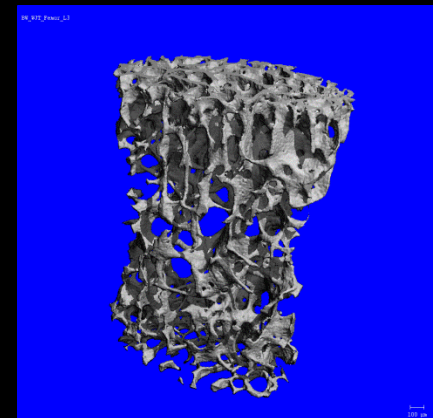
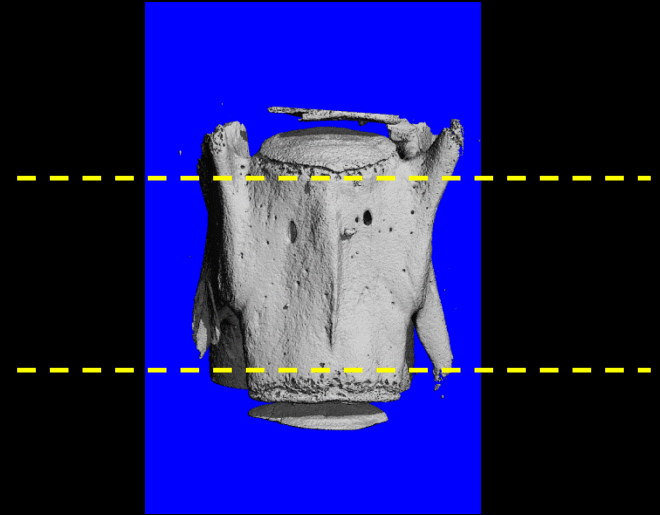
Phenotyping

- Rat cortical bone (long bone)
 - Midshaft of tibia or femur
 - 6-10 μm 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



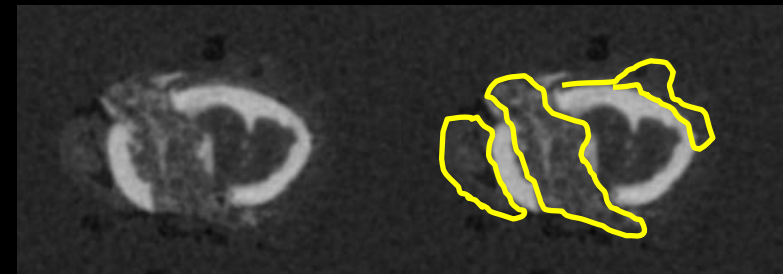
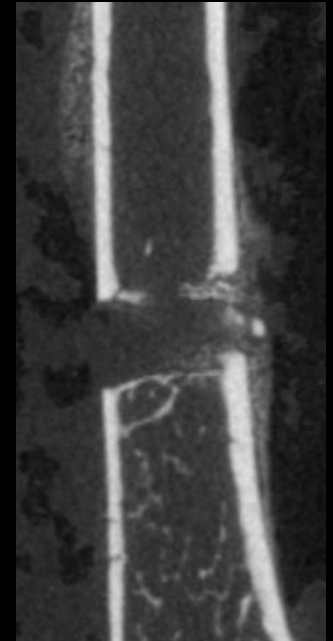
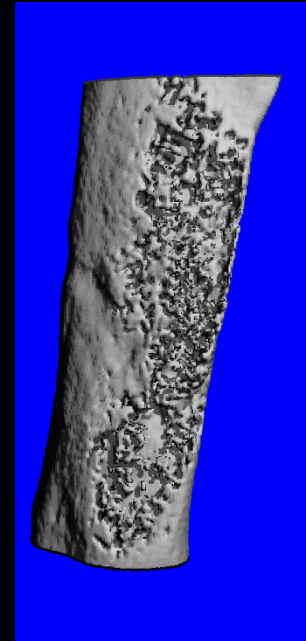
Phenotyping

- Mouse/Rat vertebral trabecular bone
 - Lumbar vertebra L1-L4
 - 6 μm (mouse) / 6-10 μm (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



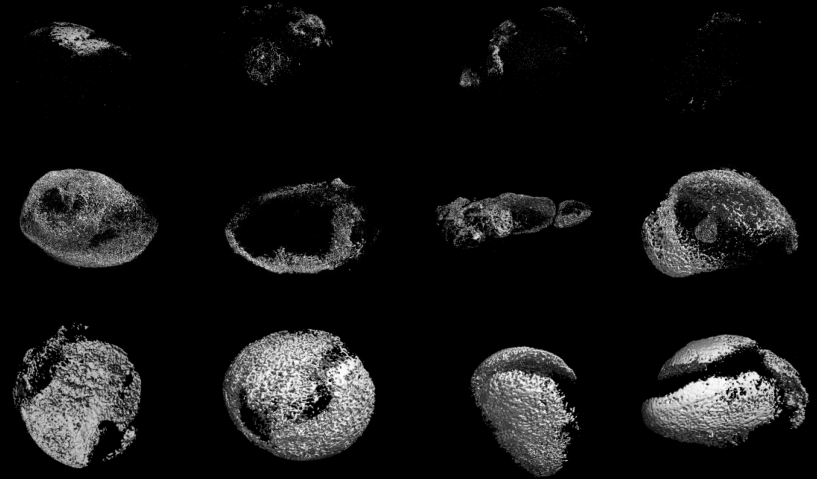
Fracture Healing

- 6-10 μm 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 μm 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

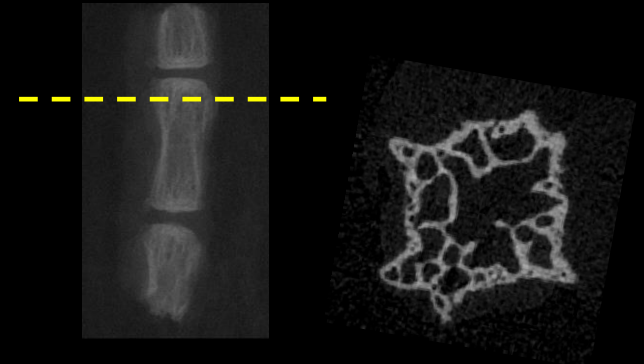


Penn

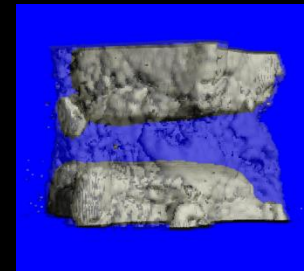
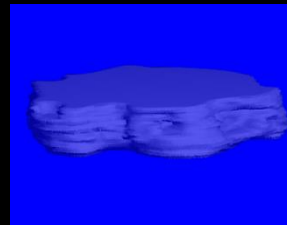
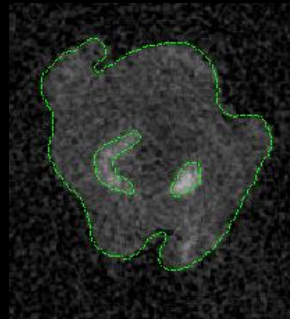
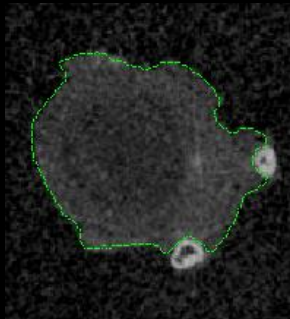
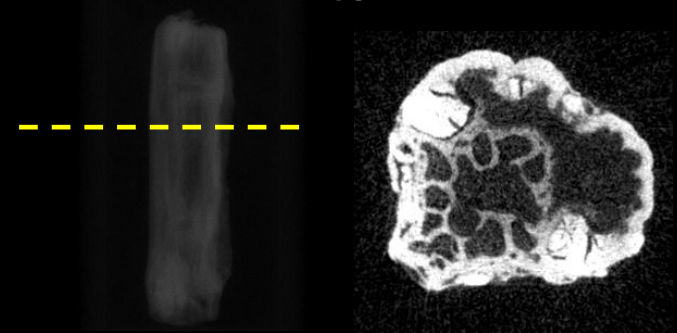
Soft Tissue μ CT Imaging

- Contrast-Enhanced μ CT Imaging
 - Mouse Intervertebral disc
 - 19 μ 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



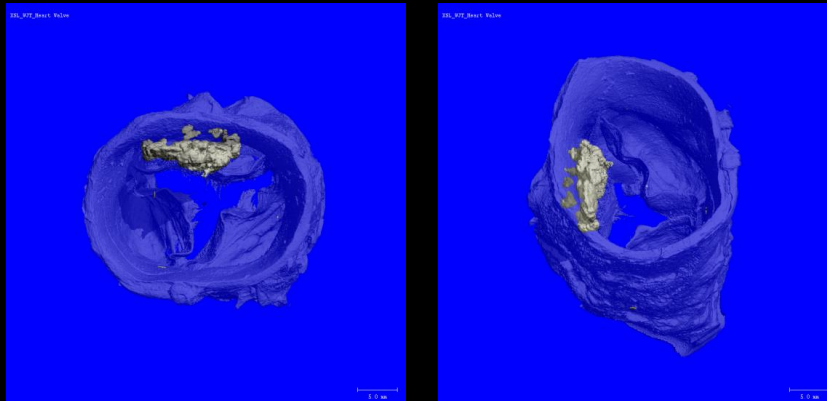
After



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Soft Tissue μ CT Imaging

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



Heart Valve



Larynx Branch



Questions?



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