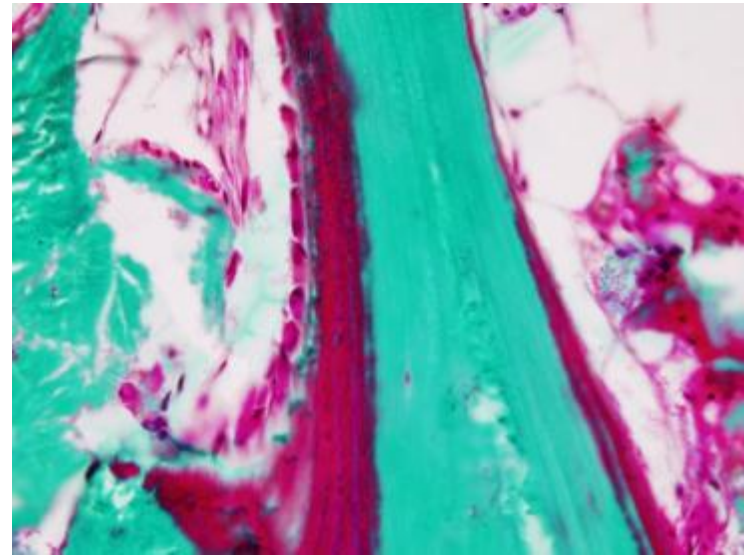
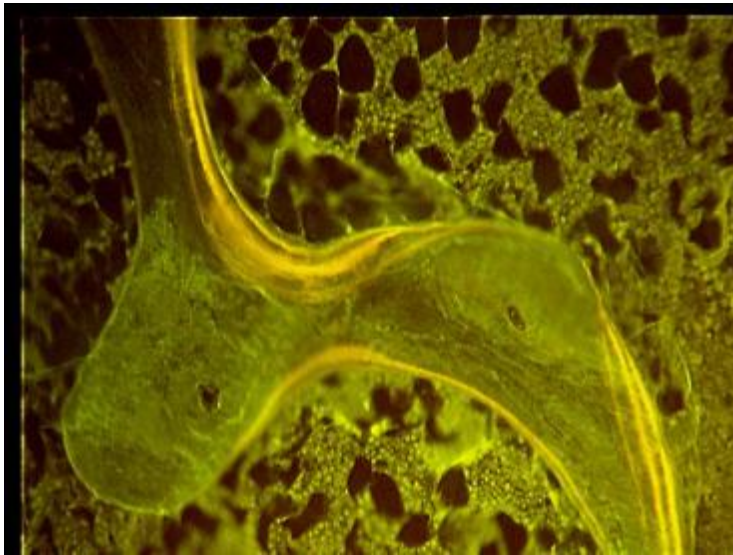


What's new in Bone Histology and Histomorphometry



Deborah Novack, MD/PhD

Director, Histology and Morphometry Core

Musculoskeletal Research Center

Division of Bone and Mineral Diseases

Departments of Medicine and Pathology

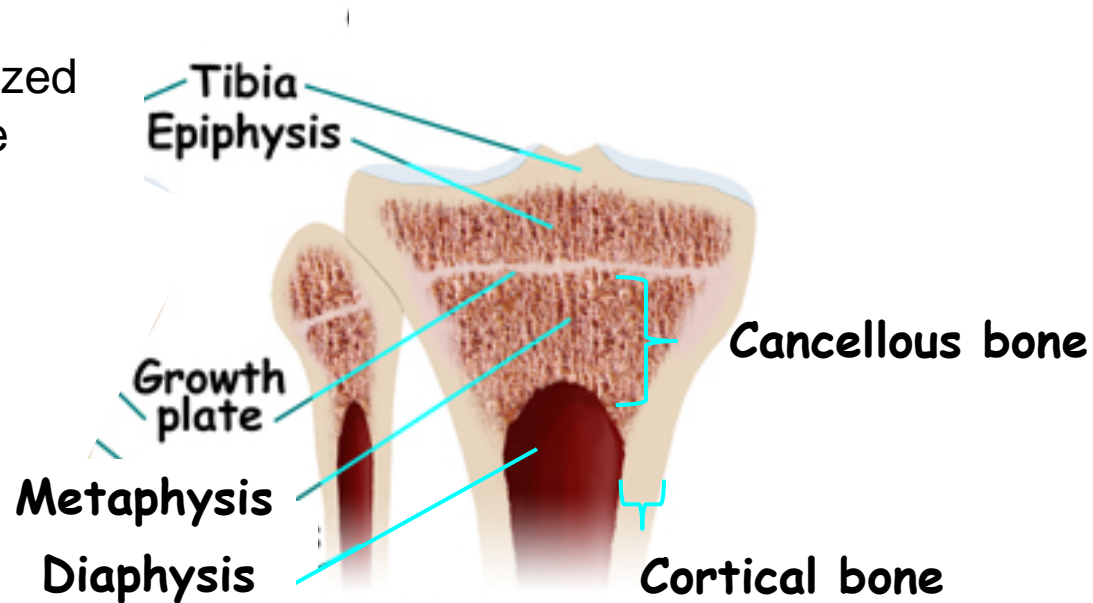
Role of Bone Histomorphometry

- histological assessment of bone phenotypes
- directly visualize bone cells in relation to tissue, at baseline or in disease models
- “quantitative” nature allows comparisons between groups even when differences are not visually “obvious”
- most useful when interpreted in the context of other data such as structural analysis (microCT, DEXA), serum markers of bone turnover, and/or disease scoring

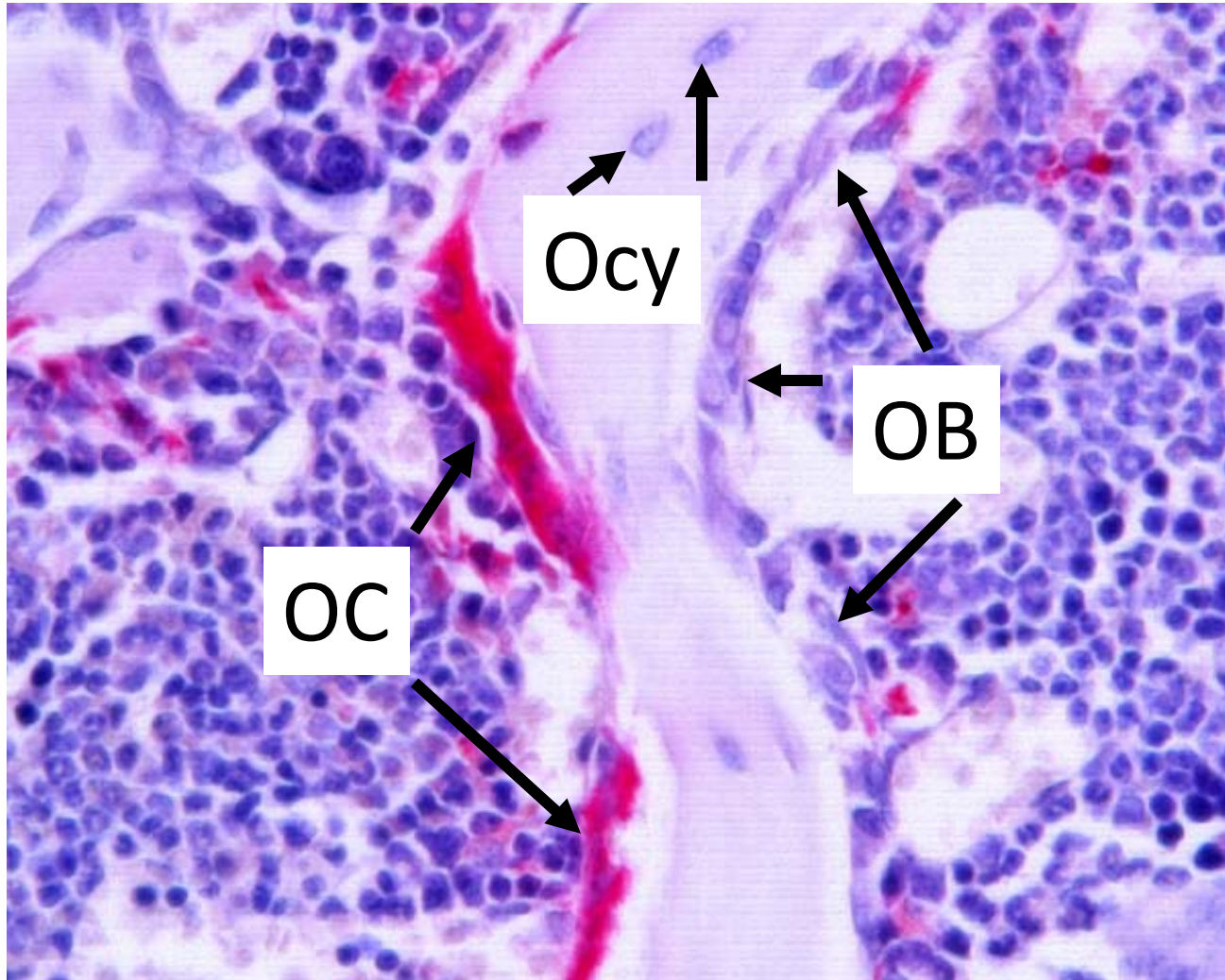
Terminology

Bone: includes mineralized and unmineralized bone matrix, but not marrow

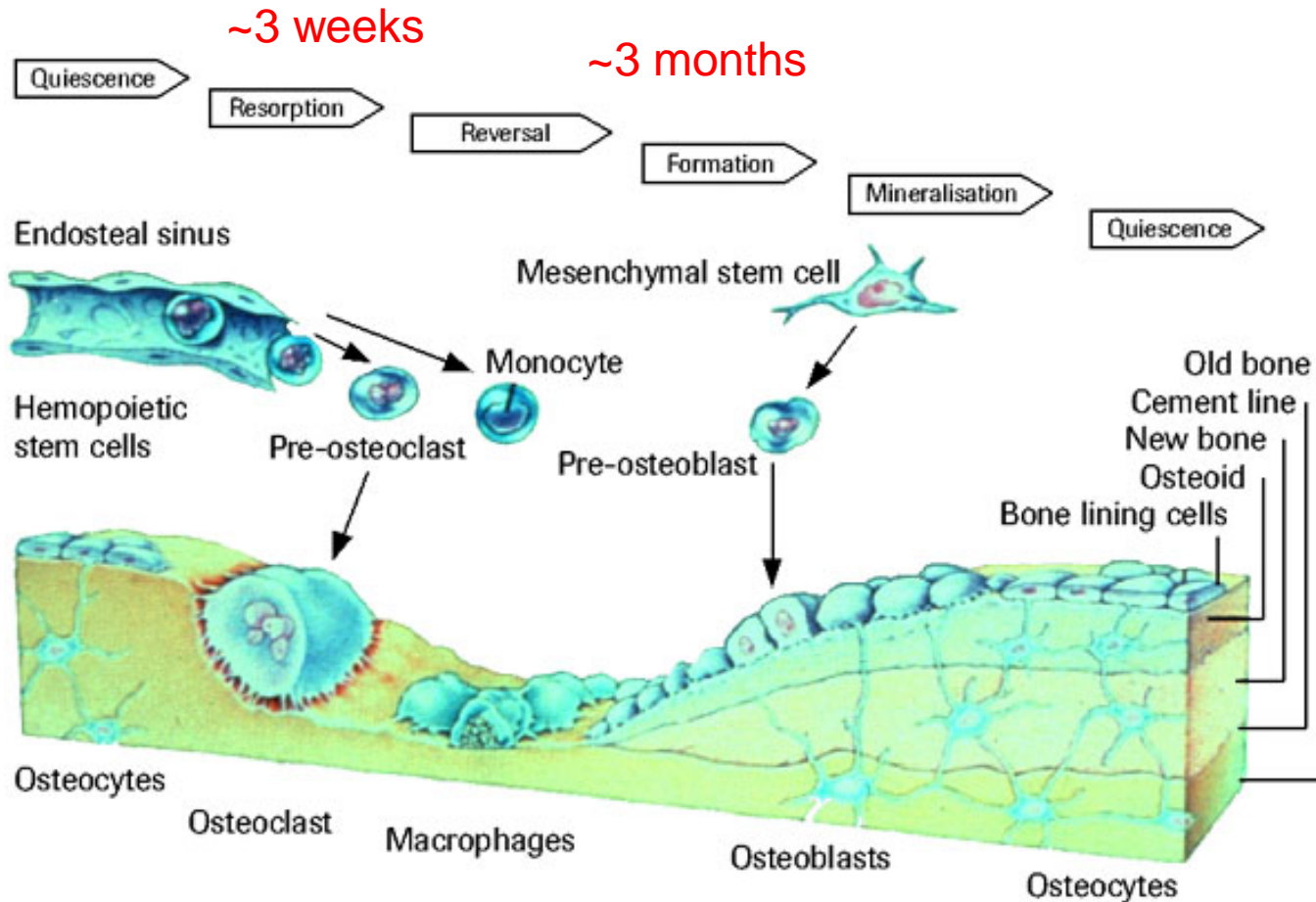
Bone tissue: includes marrow and other associated soft tissue



Trabecula/trabecular refers to an individual structural element of cancellous bone (ie a rod or plate), but **should not be used to refer to the type of bone (although we/I often do this)**



Bone Remodeling Cycle

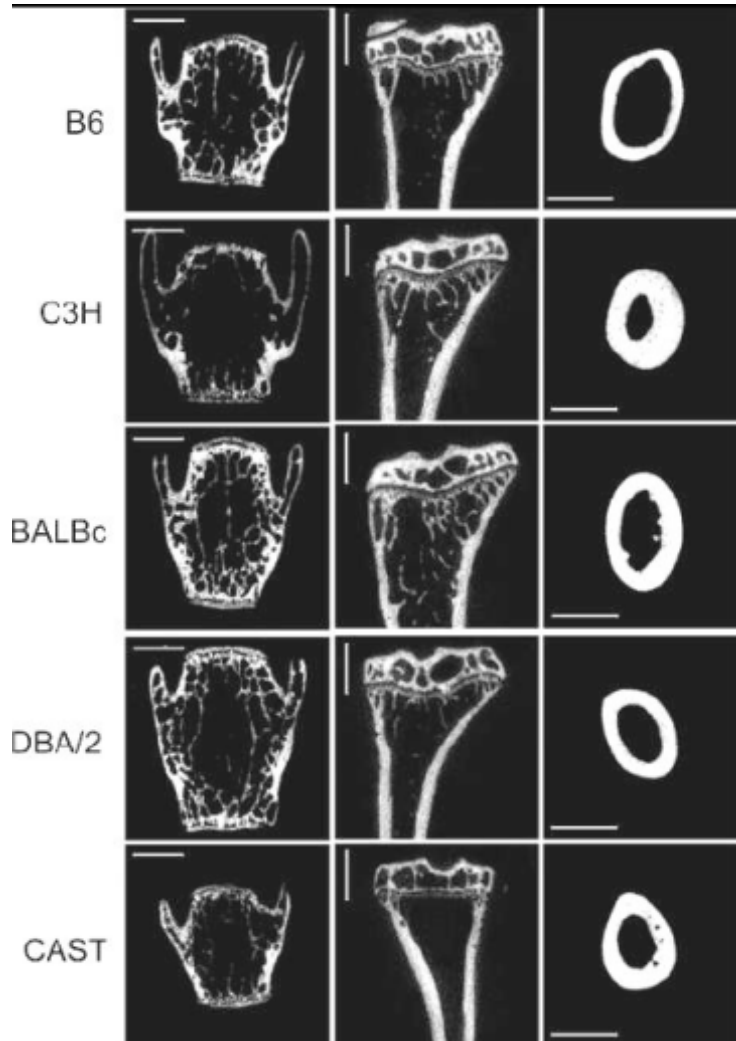


For humans

Decide what analysis you want to do
BEFORE you start your experiment!!

- There are many options for histology, and some affect how you handle the tissue when you harvest – plan ahead!
 - ✓ Fixation
 - ✓ Embedding medium
 - ✓ Orientation/plane of section
 - ✓ Type of stain

Choice of controls



Mouse strains vary a lot
C57/Bl6 may not be the “best”
since bone mass is relatively low,
but it is what we often have to use

- littermates are best
 - age/sex matched
- same strain from different suppliers are not the same in bone mass, so beware of buying controls

A quick editorial

- Although histomorphometry gives you numbers with standard deviation, p values, etc, the parameters that you measure are often not black and white, and thus are somewhat subjective
- This lack of objectivity can be especially problematic if the one measuring is biased, either consciously or subconsciously
- Therefore, you should always be **blinded**
 - have a labmate relabel your slides and keep the key
or
 - have a labmate take photos for you and code them
- The quality of your sections **DEPENDS** on your treatment of the tissue prior to processing for histology
 - Think about this as you plan your experiment, not on the day of harvest

Choice of fixation

- 10% neutral buffered formalin is the standard
 - Necessary for paraffin embedding
 - 48 hours max for TRAP stains (to see OCs), then wash in PBS and put through 30% and 50% ethanol prior to 70% for storage at 4°C
 - PFA 4% gives similar results
 - Optional for plastic/MMA embedding
 - 24 hours max if you have calcein/alizarin labels
 - Usually not used for frozen sections
- 70% ethanol
 - Samples for plastic/MMA can go directly into this
- Always use at least 10x volume of your bone!
- Excess muscle inhibits fixation

Choice of embedding method

- Plastic (methylnmethacrylate) vs. paraffin
 - plastic is used for mineralized bone, paraffin requires demineralization (eg. with EDTA)
 - bone formation (calcein labels) can only be analyzed in plastic sections
 - stains that differentiate mineralized bone from osteoid (vonKossa, trichrome) can only be done on plastics
 - BUT, plastic takes longer and is more expensive

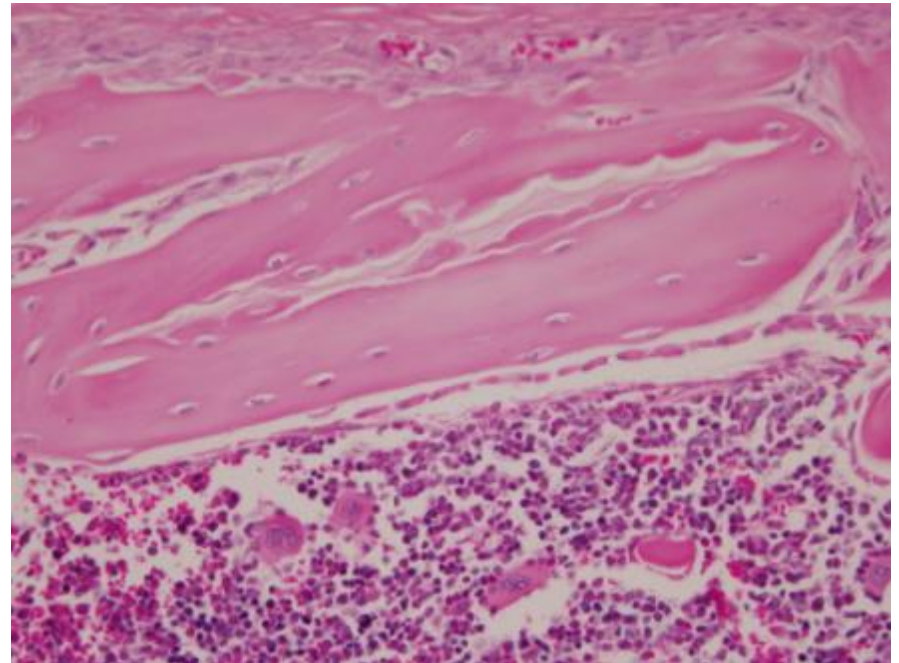
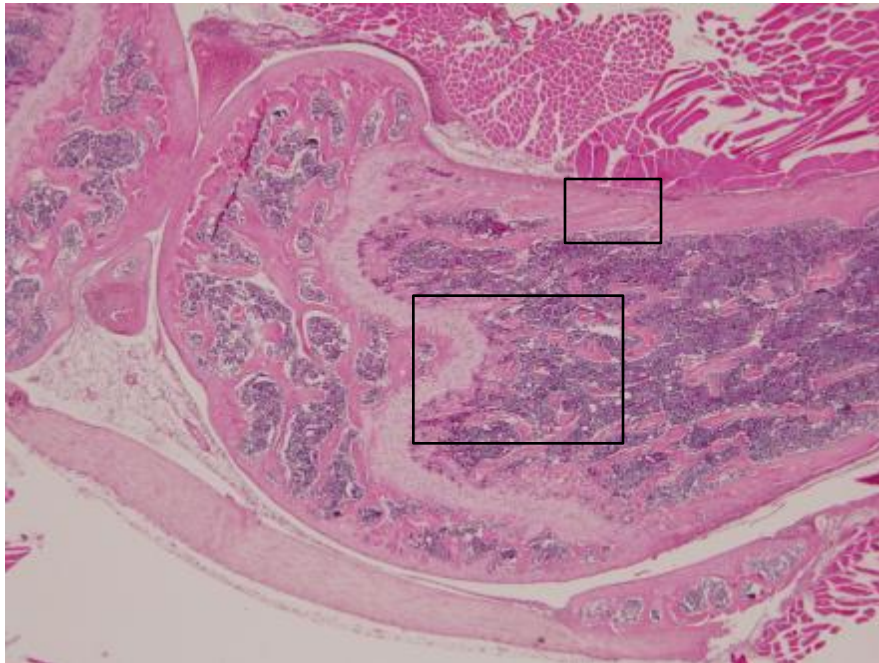
Frozen section options (see later) are not used for standard histomorphometry

Tissue orientation

- Since histology gives you a 2-D image, you need to think about exactly what you are trying to capture
- If you need a very particular plane of section, it is useful to provide a low power image of some sort (can even be a uCT slice) to help guide the sectioning

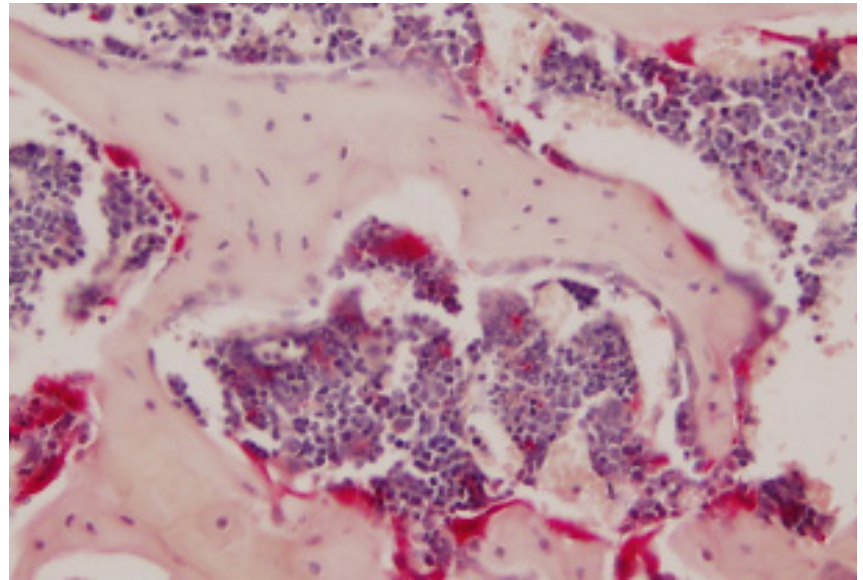
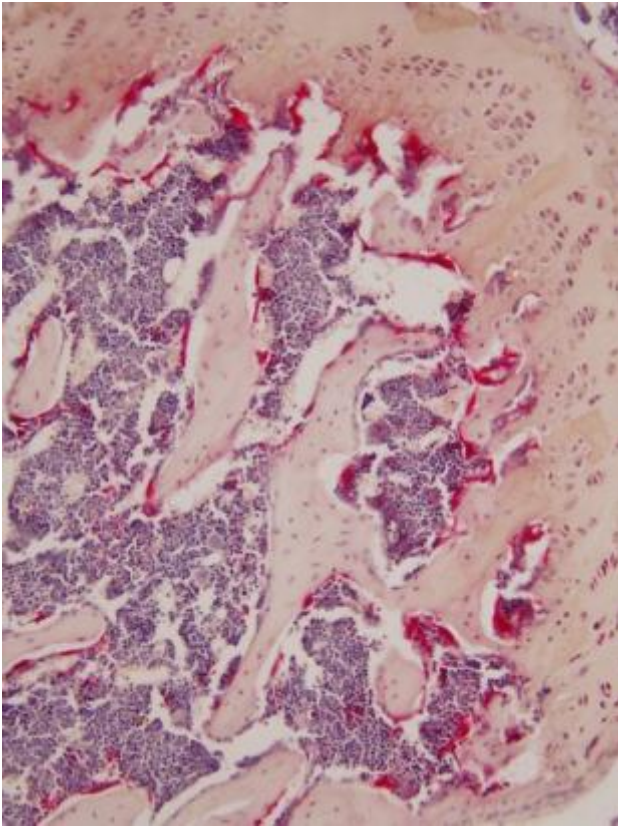
Choice of stain – H&E

- The “bread and butter” of standard histology
- good for seeing bone vs cartilage (growth plate) and seeing OBs, but not OCs, also good to see tumor or inflammation



Choice of stain – TRAP

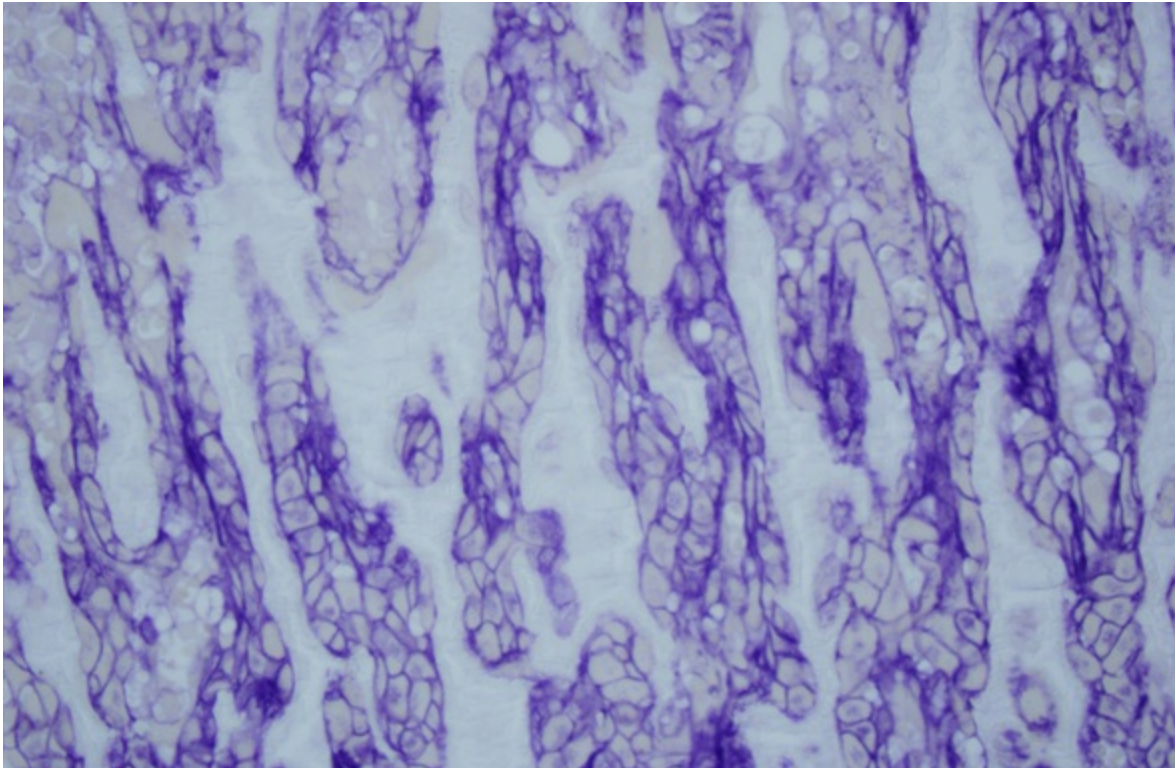
- highlights OCs, still OK for bone vs cartilage, OBs usually identifiable, often used for histomorph
- Works on paraffin or plastic



OCs are often defined as TRAP+ cells on bone surface with > 1 nucleus, but sometimes they have 1 or none in a section. ASBMR says to state criteria you use.

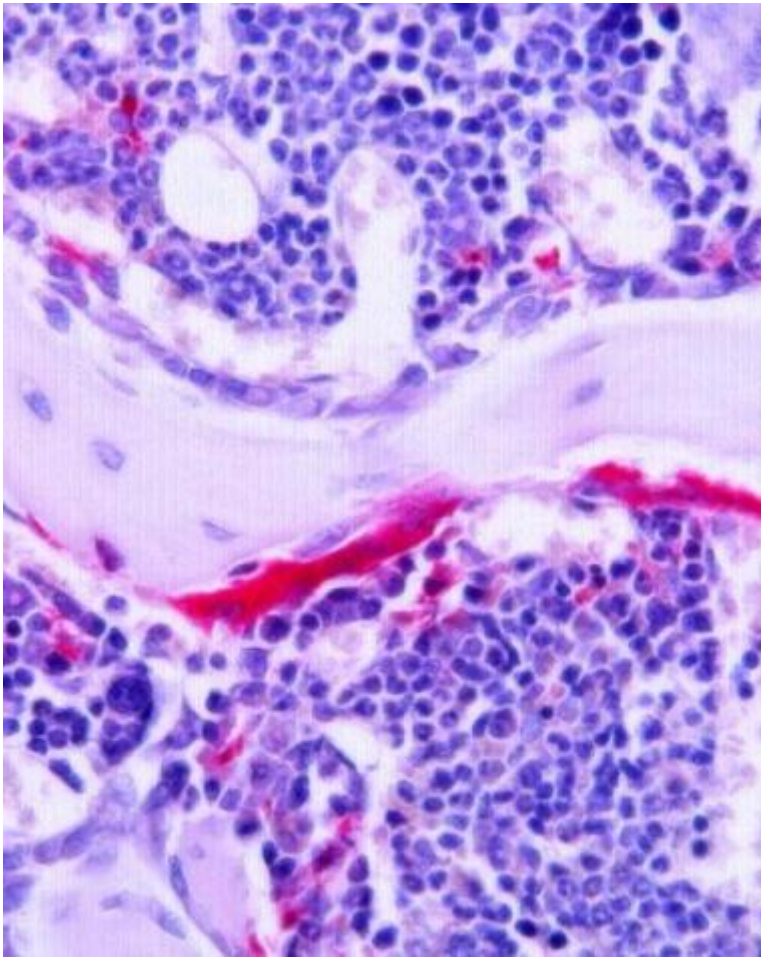
Choice of stain – Alkaline phosphatase

- Highlights OBs, can't see OCs, decreased by formalin fixation



Not currently offered routinely in our core; being developed for MMA and frozen sections

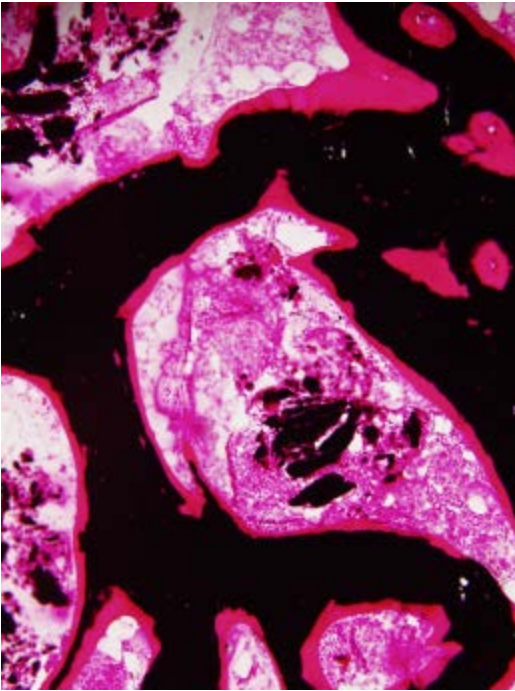
A word on counting osteoblasts



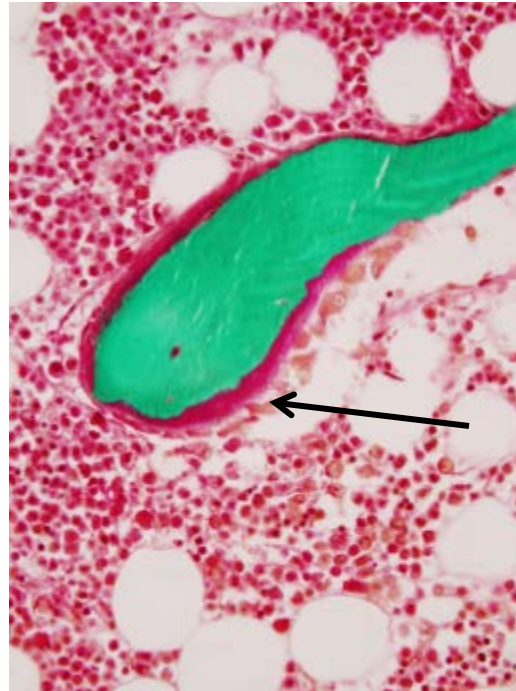
- ASBMR taskforce says that term Ob should be restricted to active cells, and not flat lining cells (which are also of Ob lineage)
- With or without an AP stain, Obs should be cuboidal in shape
- Counting Obs often feels very subjective – try to be consistent
- Back up counts with calcein labels and/or serum markers

Identifying osteoid

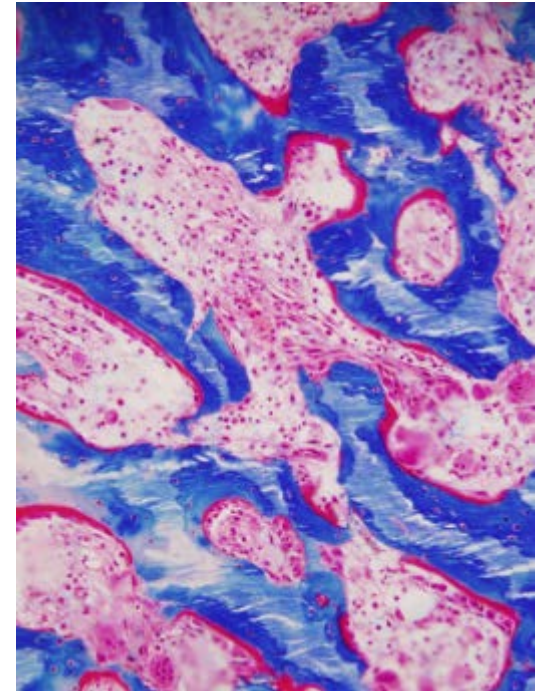
- vonKossa – shows osteoid vs bone, not good for cellular detail
- trichrome (Masson or Goldner) – shows osteoid, and better for cellular detail



vonKossa



Goldner trichrome

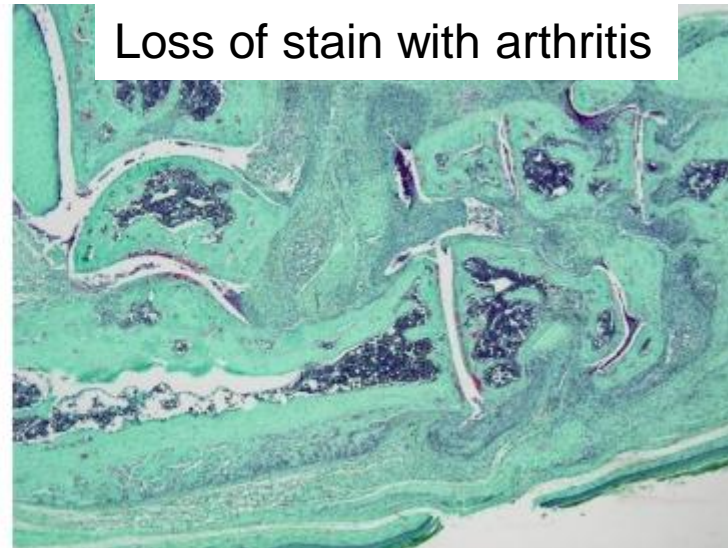
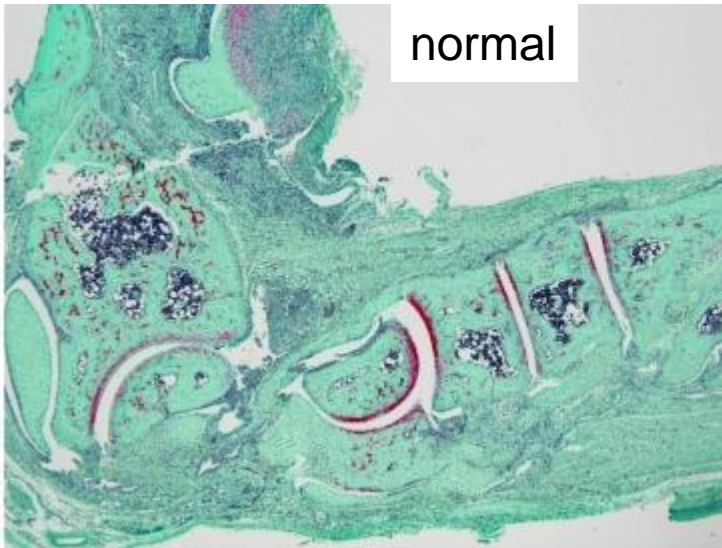


Masson trichrome

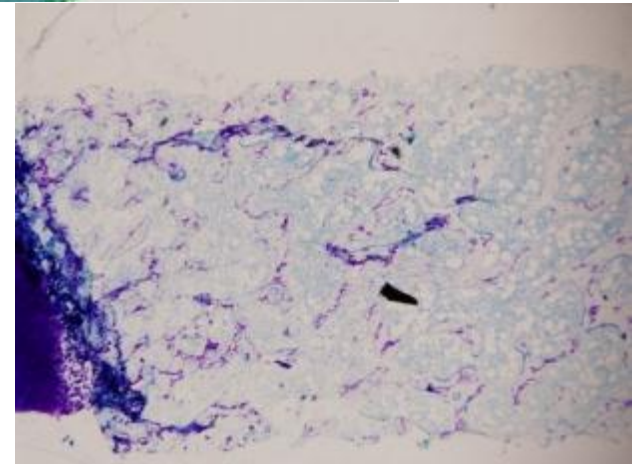
Identifying cartilage

- safranin O – red with methyl green counterstain
- toluidine blue – deep blue/purple

safranin O



toluidine
blue



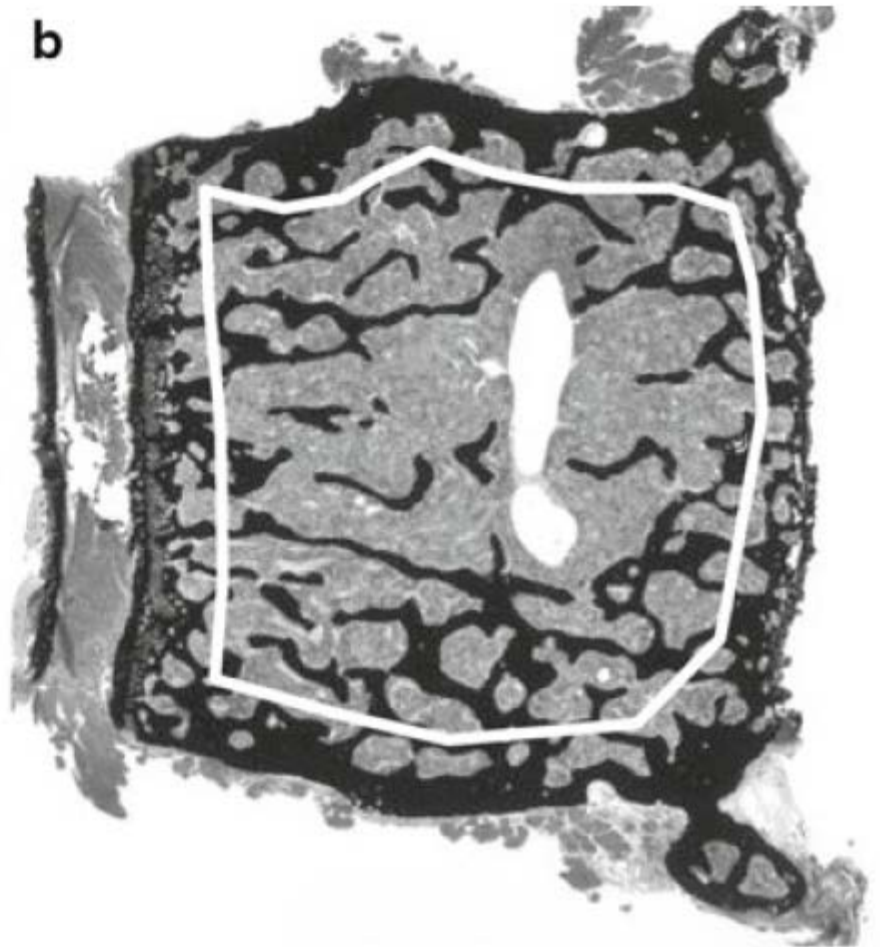
A brief advertisement...

- This histology core (aka Core C) provides a number of services including various embedding and staining options
- Crystal is not only skilled with the various techniques, but can be an important resource during both the planning and troubleshooting phases of your projects
 - Understanding the histochemical basis for the stains, she can help choose fixative to work with a stain, or the right embedding procedure
 - Looking in the microscope, she can help determine if problems are due to tissue preparation or staining

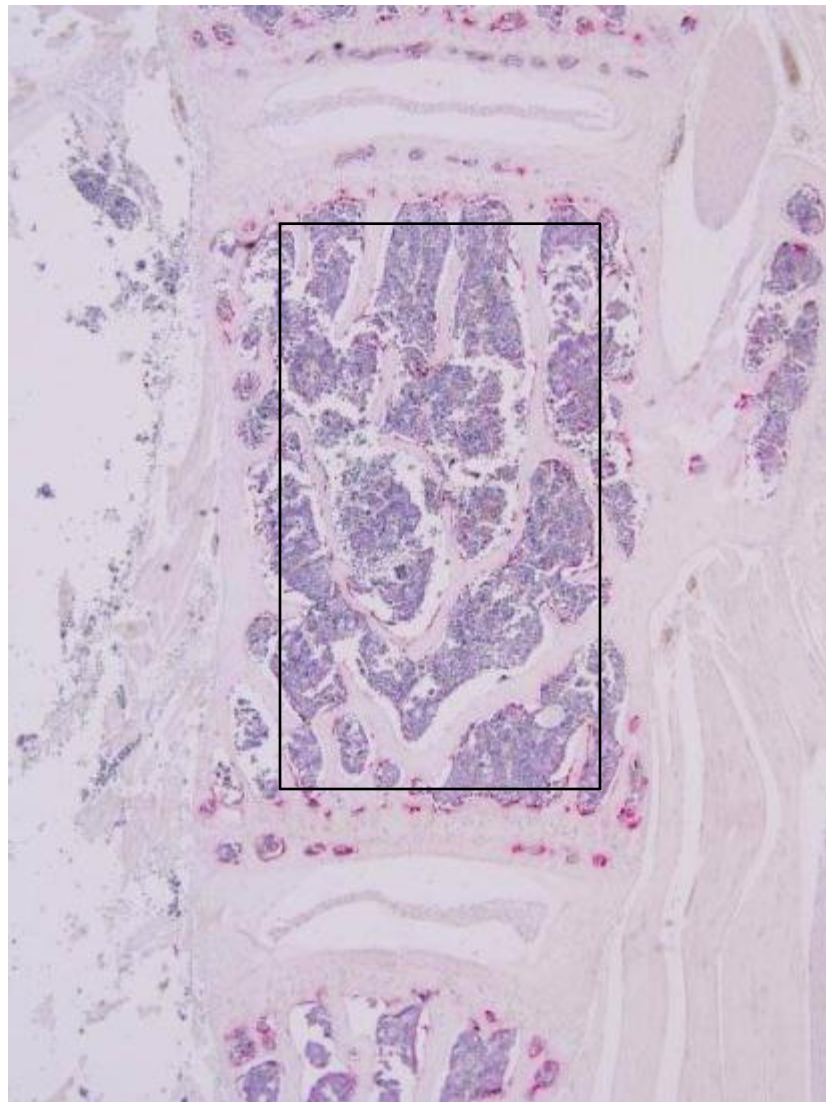
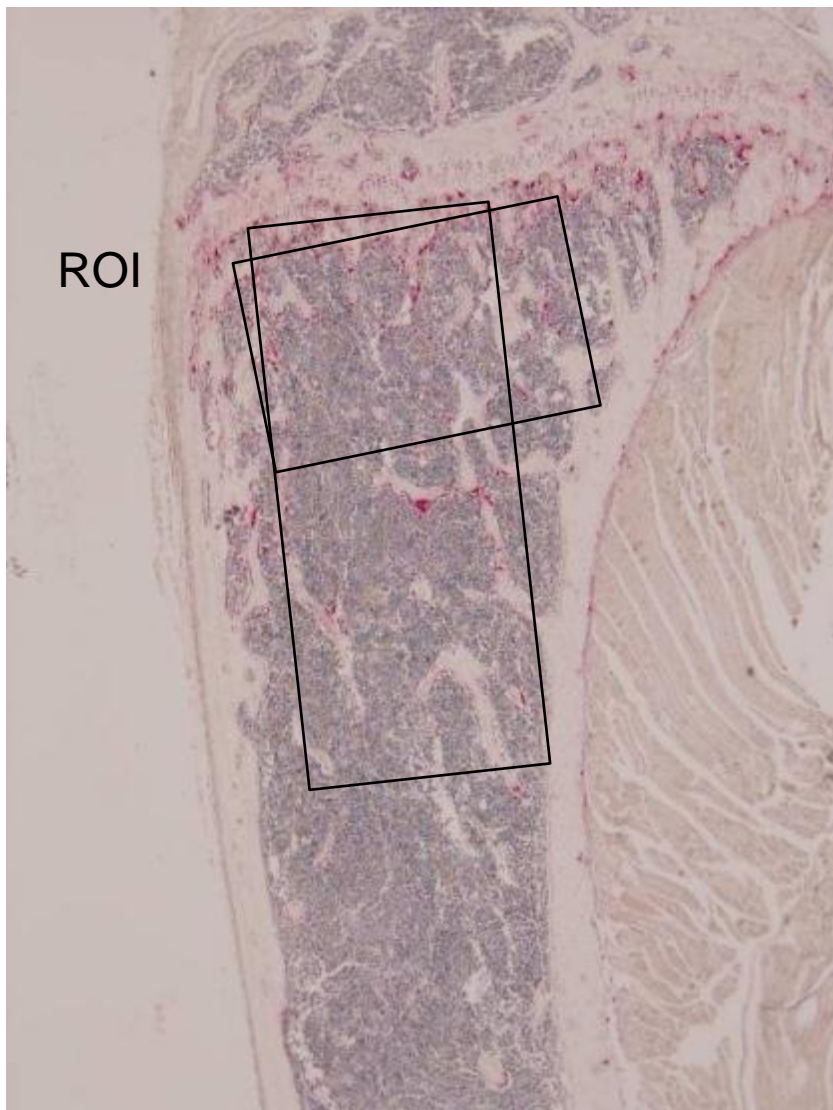
So I have stained slides, now what?

- Prior to beginning analysis with BioQuant (histomorphometry), browse your sections
 - Are they all adequate or do you need some recuts?
 - Which regions do you need to analyze?
 - Which parameters are most important?
 - Are you defining a basal phenotype or looking at a disease model?
- Keep in mind that the ASBMR documents were designed primarily for basal phenotypes and osteoporosis, but the BioQuant software can accommodate many other situations

Define your region of interest



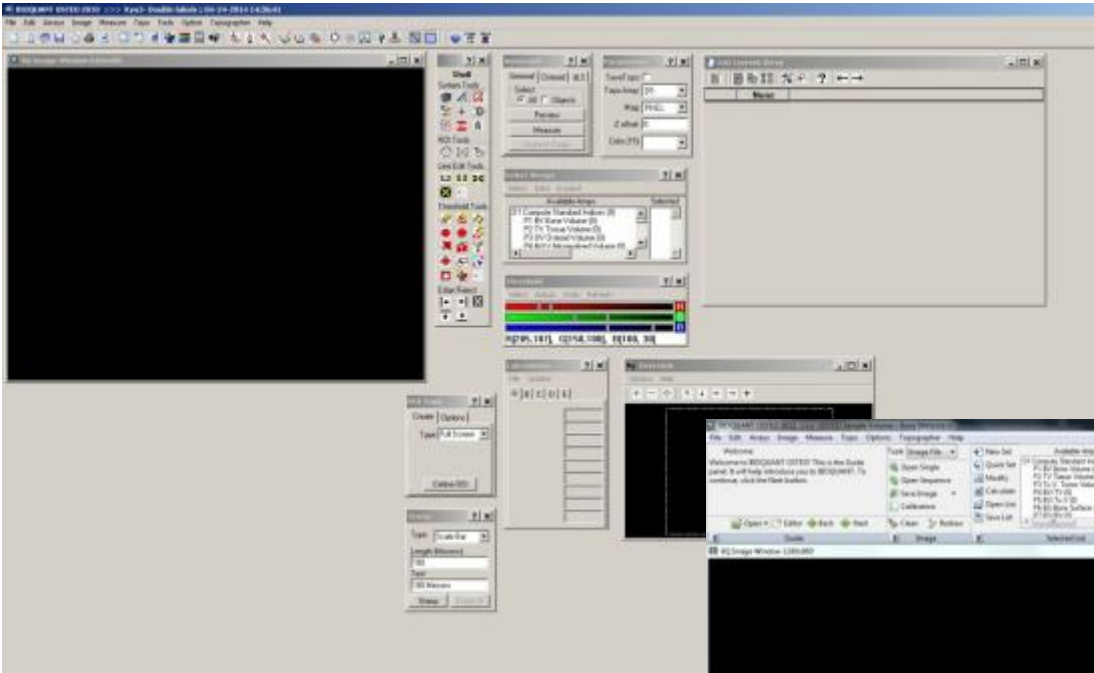
Define your region of interest



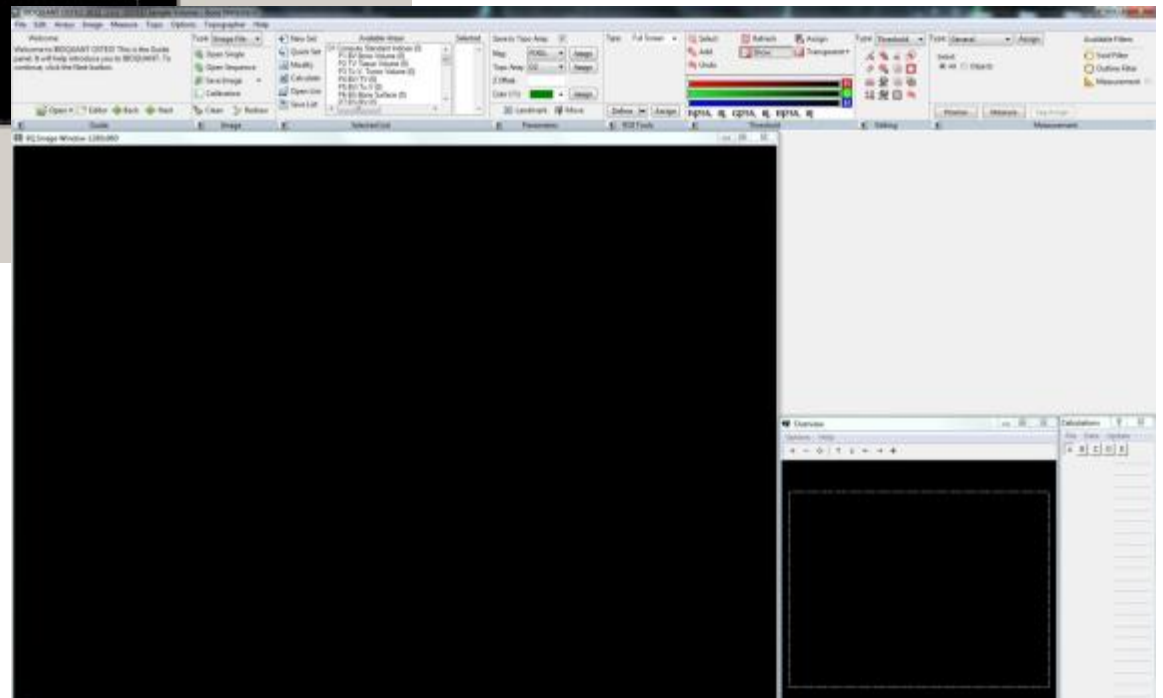
A brief digression

-- New Bioquant upgrade

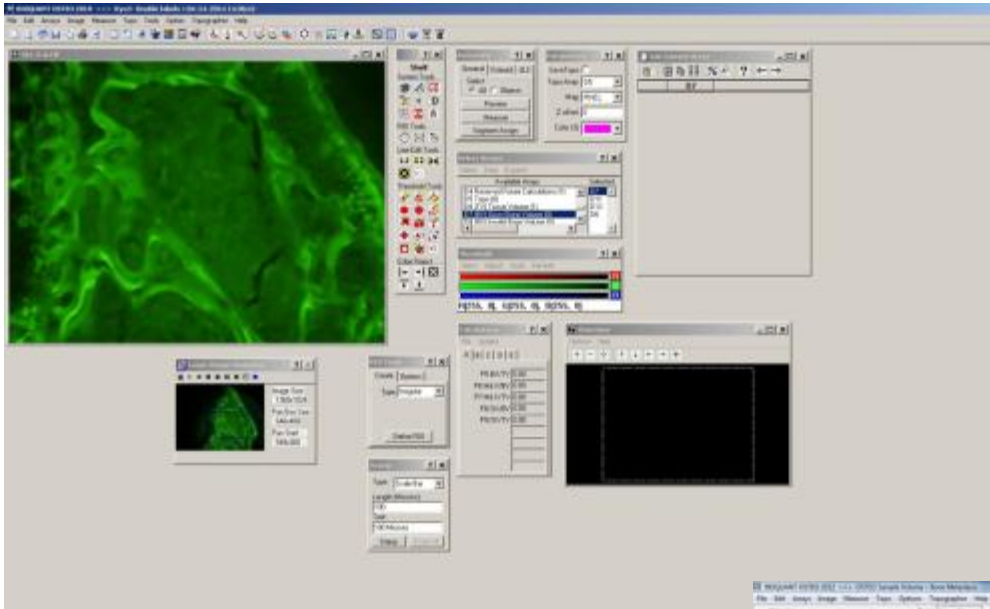
Old version



New version with task bar at top

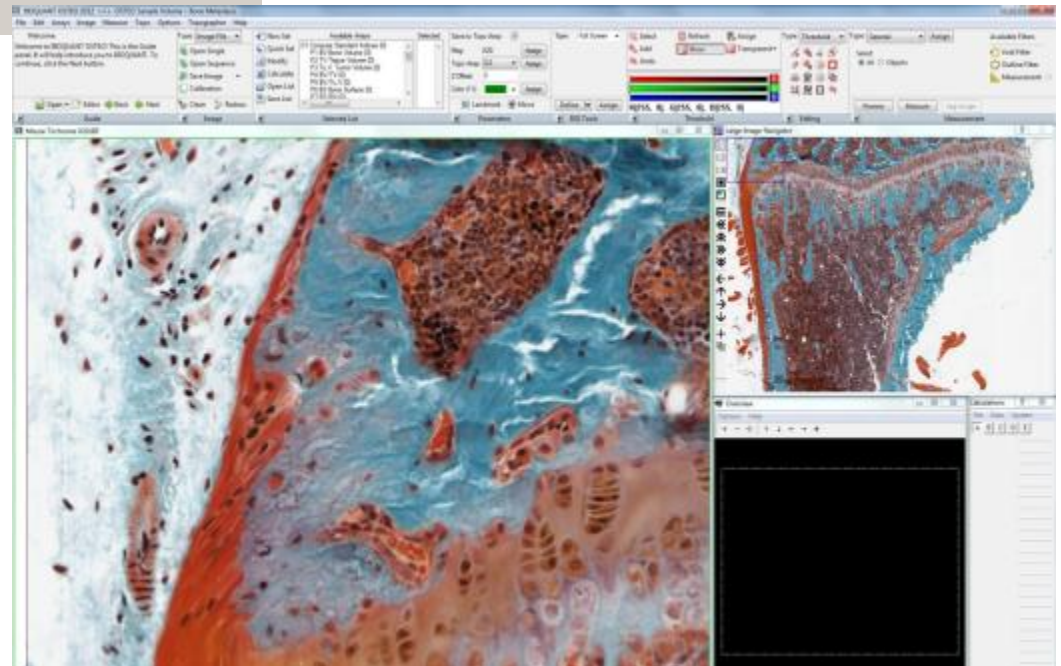


Old version: Image Window only 640x480 pixels.



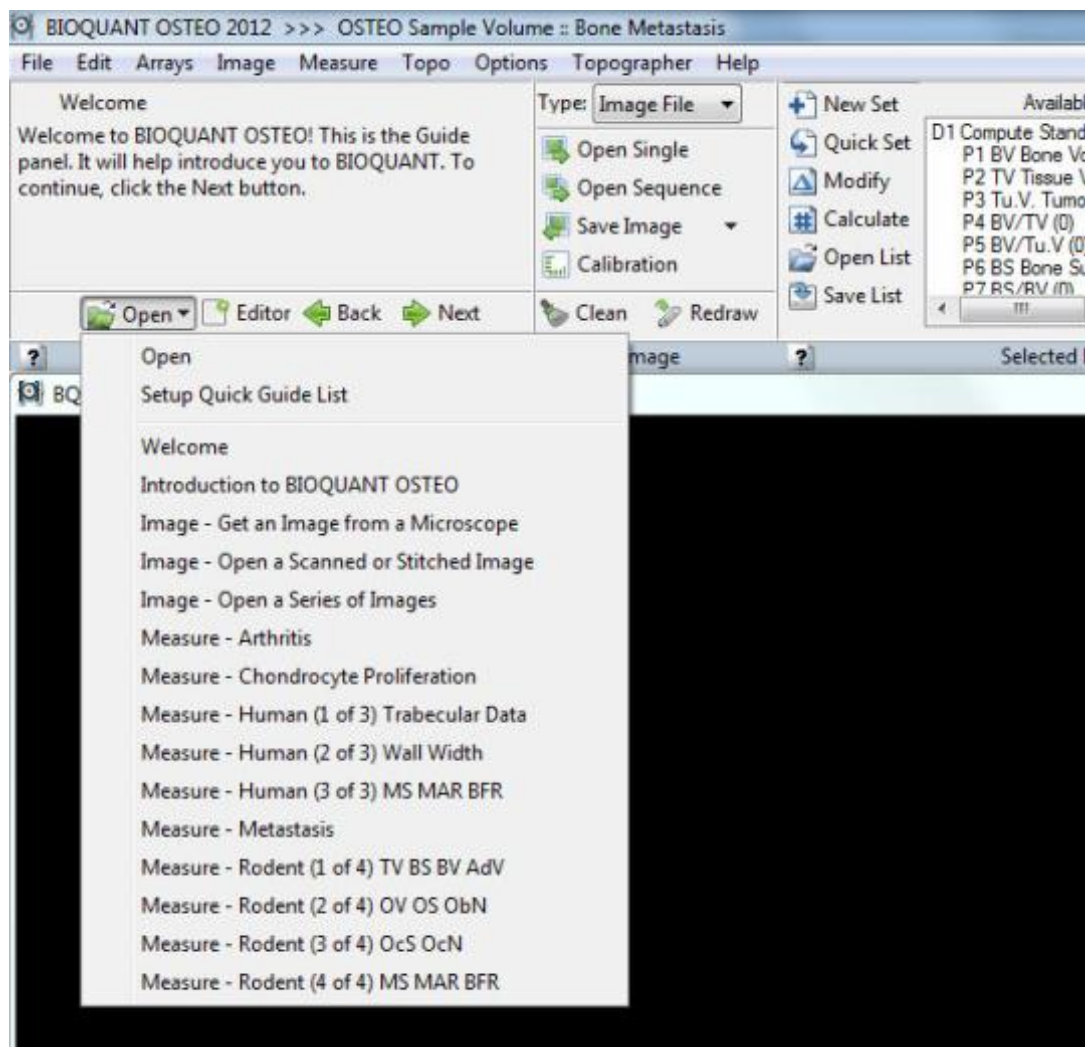
Step-up magnification (keyboard Z button) to measure features directly in Image Window. No need to move around smaller ROI box to cover your overall ROI.

New version:
Image Window up to 1280x960
pixels.



4. Detailed tutorial is provided in 2012 version

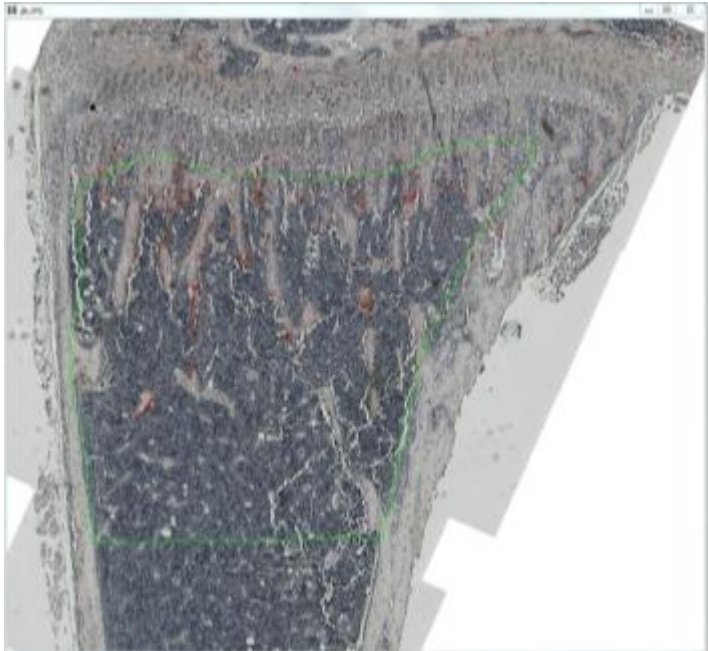
Tutorials to walk you through most the common analyses



Additional upgrades in new Bioquant

- Assign buttons to fix the unique settings in current analysis, such as magnification, ROI type, manual threshold, etc. No need to reset between samples.
- One array (L3) finishes both BV and BS measurement.
- Automatic thresholding still only useful with a high contrast stain like trichrome, not TRAP or H&E

Unmagnified photo in Image Window



2x magnification in Image Window



Standard nomenclature

- ASBMR taskforce has a list of individual terms
 - All except BMU (basic multicellular unit) are 1 or 2 letters, the first being capitalized and second lower case
- Standard format = Source – Measurement/Referent; period between 2 terms if double letter
 - Eg. Cn-N.Oc/BS means cancellous Oc number per bone surface
 - If same source (cancellous or cortical) is used repeatedly, don't need to keep indicating
 - The referent (TV, BV, BS, or OS most commonly) is very important and should never be omitted

Table 2. Sources and Referents in Bone Histomorphometry

Sources		Referents	
Name	Abb.	Name	Abb.
Total core	Tt	Bone surface	BS
Cortical bone tissue	Ct	Bone volume	BV
Cancellous bone tissue	Cn	Tissue volume	TV
Endocortical surface	Ec	Core volume	CV
Periosteal surface	Ps	Osteoid surface	OS
Transitional zone	Tr.Z	Bone interface	BI
Diaphyseal bone	Dp	Eroded surface	ES
Metaphyseal bone	Mp	Mineralized surface	Md.S
Epiphyseal bone	Ep	Osteoblast surface	Ob.S
Medullary bone	Me	Osteoclast surface	Oc.S

Abb. = abbreviation.

Those listed will cover most situations in both human and nonhuman studies, but neither list is exhaustive. Combinations of source terms may be needed, such as Dp.Ec for diaphyseal bone, endocortical surface.

Slides are 2-dimensional but most reported histomorphometric parameters are 3-D

- The goal of histomorphometry is to understand 3-dimensional reality
 - Stereology and mathematics provide approaches to convert data to 3-D, but most rely on assumptions such as isotropy which are not strictly true for bone
 - ASBMR recommends that 3-D terminology be used for volume, surface, and thickness, although numbers of cells remains in 2-D (ie. Referent is a surface, not a volume)
 - 2-D perimeter/area ratios are converted to 3-D surface/volume by multiplying by $4/\pi$ or 1.2
 - The Bioquant software does this for you

Table 3. Primary Measurements in Bone Histomorphometry

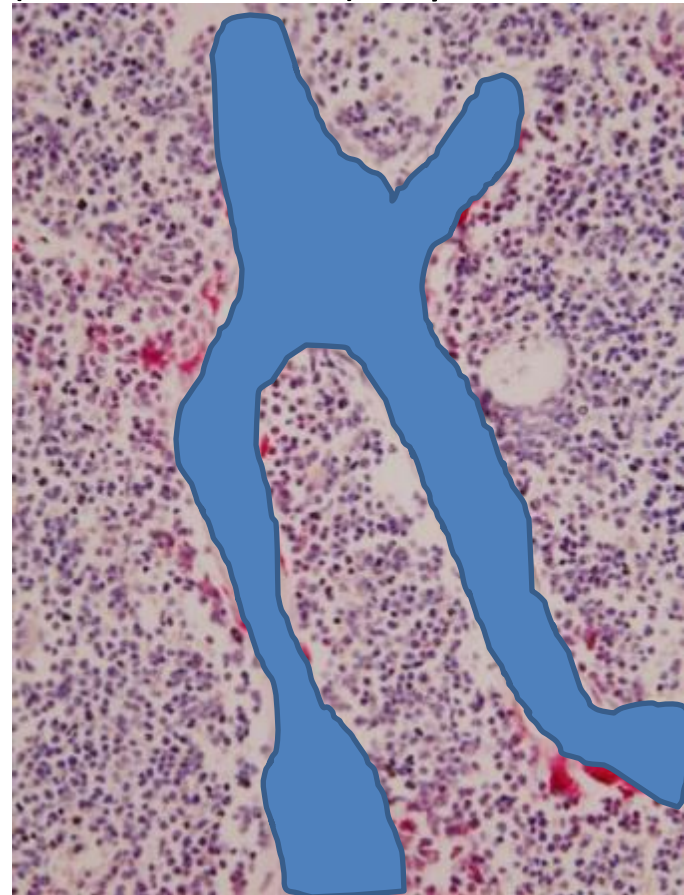
Type of measurement	Name of measurement	Abbreviations		Type of measurement	Name of measurement	Abbreviations	
		3D	2D			3D	2D
Area	Bone volume ^a	BV	B.Ar	Distance ^k	Cortical thickness ^l	Ct.Th	Ct.Wi
	Osteoid volume	OV	O.Ar		Wall thickness	W.Th	W.Wi
	Mineralized volume	Md.V	Md.Ar		Mineralized thickness	Md.Th	Md.Wi
	Void volume	Vd.V	Vd.Ar		Osteoid thickness	O.Th	O.Wi
	Marrow volume	Ma.V	Ma.Ar		Label thickness	L.Th	L.Wi
	Fibrosis volume	Fb.V	Fb.Ar		Trabecular thickness ^m	Tb.Th	Tb.Wi
	Canal volume ^b	Ca.V	Ca.Ar		Interstitial thickness	It.Th	It.Wi
	Cell volume ^{b,c}	Ce.V	Ce.Ar		Trabecular diameter ⁿ	Tb.Dm	— ^o
	Cytoplasmic volume ^{b,d}	Cy.V	Cy.Ar		Canal radius	Ca.Rd	— ^o
	Nuclear volume ^{b,d}	Nc.V	Nc.Ar		Cell height ^c	Ce.Ht	— ^o
	Length	Bone interface ^e	BI		B.Bd	Number ^p	Nuclear height ^d
Bone surface ^f		BS	B.Pm	Erosion depth	E.De		— ^o
Osteoid surface		OS	O.Pm	Osteoblast number	—		N.Ob
Eroded surface		ES	E.Pm	Osteoclast number	—		N.Oc
Quiescent surface ^g		QS	Q.Pm	Osteocyte number	—		N.Ot
Mineralized surface ^h		Md.S	Md.Pm	Adipocyte number	—		N.Ad
Osteoblast surface		Ob.S	Ob.Pm	Nuclear number ^d	—		N.Nc
Single-labeled surface ⁱ		sLS	sL.Pm	Canal number	—		N.Ca
Double-labeled surface ⁱ		dLS	dL.Pm	Seam number	—		N.Sm
Osteoclast surface		Oc.S	Oc.Pm	Erosion number	—		NE
Reversal surface ^j		Rv.S	Rv.Pm	Profile number	—		N.Pf

In rodent models, additional parameters such as thickness of growth plate or articular cartilage might be useful, and the ASBMR “lexicon” provides appropriate symbols

Things you measure directly

- Area (volume) – outlines a closed space
 - bone volume (BV) = amount of bone in your ROI
 - tissue volume (TV) = amount of tissue (bone+ marrow) in your ROI
 - osteoid volume (OV)
- Length - lines
 - bone surface (BS)
 - osteoblast surface (Ob.S)
 - single labeled surface (sLS)
- Distance – between 2 lines
 - osteoid thickness (O.Th)
 - interlabel distance (Ir.L.Th)
- Number - dots
 - osteoblast number (N.Ob)
 - osteoclast number (N.Oc)

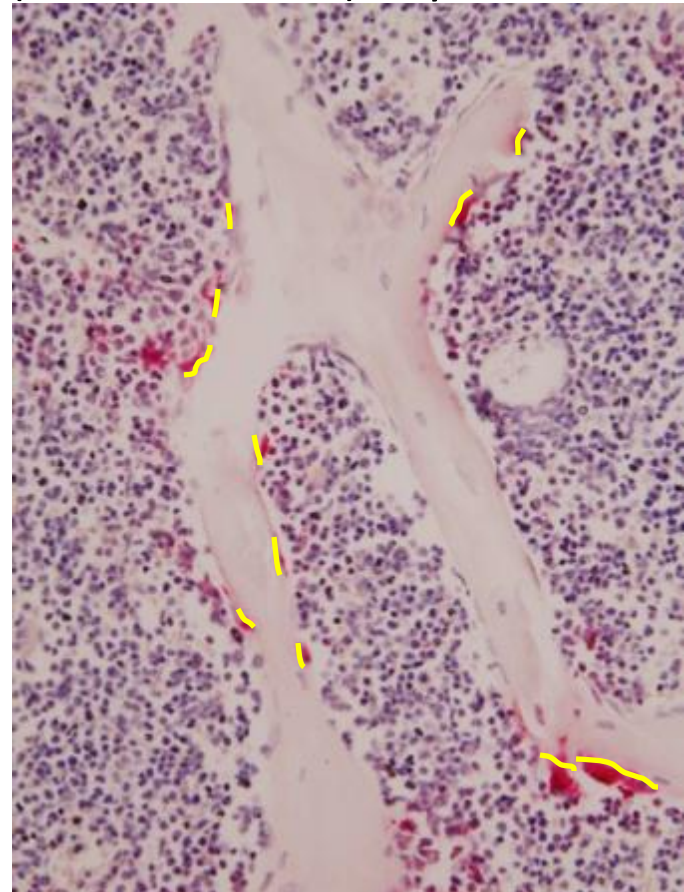
The so-called primary indices are then calculated from these – eg BV/TV or $N.Ob/BS$



Things you measure directly

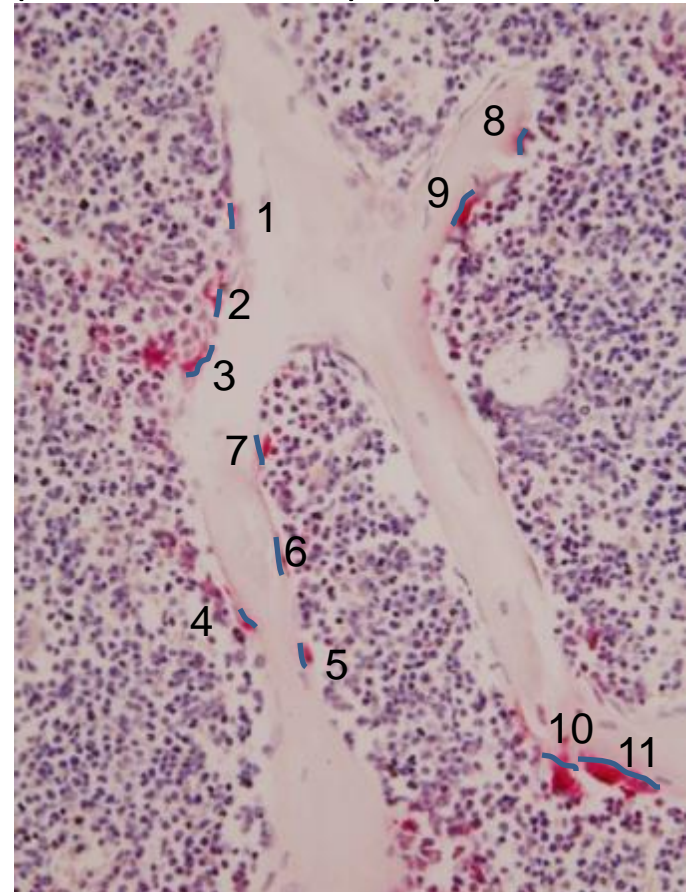
- Area (volume) – outlines a closed space
 - bone volume (BV) = amount of bone in your ROI
 - tissue volume (TV) = amount of tissue (bone+ marrow) in your ROI
 - osteoid volume (OV)
- Length - lines
 - bone surface (BS)
 - osteoclast surface (Oc.S)
 - single labeled surface (sLS)
- Distance – between 2 lines
 - osteoid thickness (O.Th)
 - interlabel distance (Ir.L.Th)
- Number - dots
 - osteoblast number (N.Ob)
 - osteoclast number (N.Oc)

The so-called primary indices are then calculated from these – eg BV/TV or $N.Ob/BS$



Things you measure directly

- Area (volume) – outlines a closed space
 - bone volume (BV) = amount of bone in your ROI
 - tissue volume (TV) = amount of tissue (bone+ marrow) in your ROI
 - osteoid volume (OV)
- Length - lines
 - bone surface (BS)
 - osteoclast surface (Oc.S)
 - single labeled surface (sLS)
- Distance – between 2 lines
 - osteoid thickness (O.Th)
 - interlabel distance (Ir.L.Th)
- Number - dots
 - osteoblast number (N.Ob)
 - osteoclast number (N.Oc)



The so-called primary indices are then calculated from these – eg BV/TV or N.Ob/BS

A note on eroded surface (ES)

- eroded surface is the “new” surface of the bone generated by an active OC, and appears “crenated” or “lacunar”



- ES can be considered Oc+ if a cell is present, or Oc-

In mice, however, it is often hard to see nice pits since the OCs seem to sit on a flat bone surface. I consider ES in mice to be very subjective, compared to Oc.S (which still involves some judgment calls).

Derived structural indices

(things that get calculated from your primary measurements based on some assumptions)

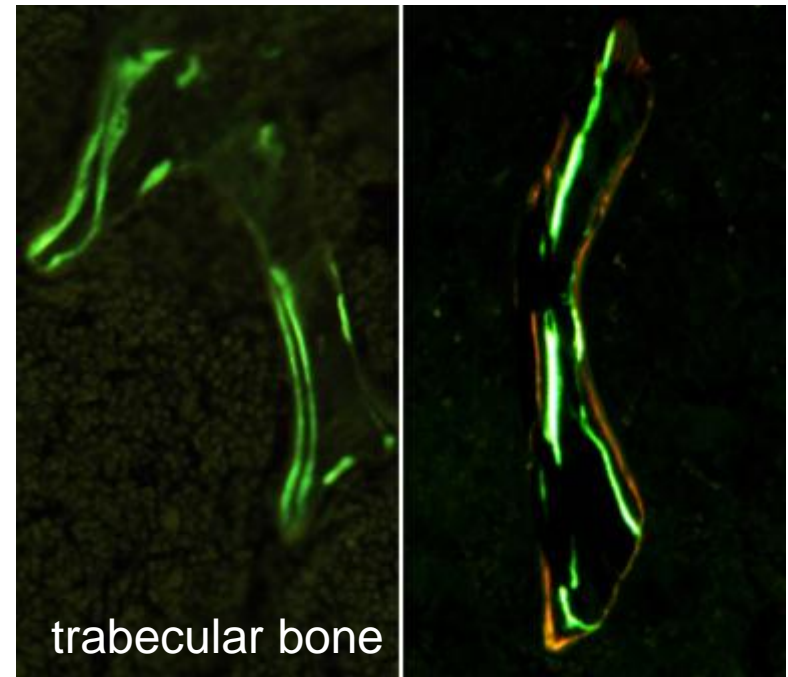
– trabecular number $Tb.N = (BV/TV)/Tb.Th$

– trabecular separation $Tb.Sp = (1/Tb.N) \cdot Tb.Th$

- These, as well as BV/TV , are better assessed by microCT
- However, in order to get values for cell counts and surfaces, you always have to measure the bone area and bone perimeter on your sections
 - Which then get converted to BV and BS

Dynamic histomorphometry

- measurement of bone formation parameters
- mice are given 2 doses of label several days apart, by IP injection
- calcein (green) and/or alizarin red (red) incorporate into newly calcifying bone
- interval depends on age of mice
 - 2-4 week mice, 3-4d
 - 6-12 week mice, 5-7d
 - older mice, 7-10d
- easier to measure in calvaria or endocortical surface than trabecular bone
 - Always should indicate where measurement was taken
 - Sometimes get different results from different sites



Derived kinetic indices

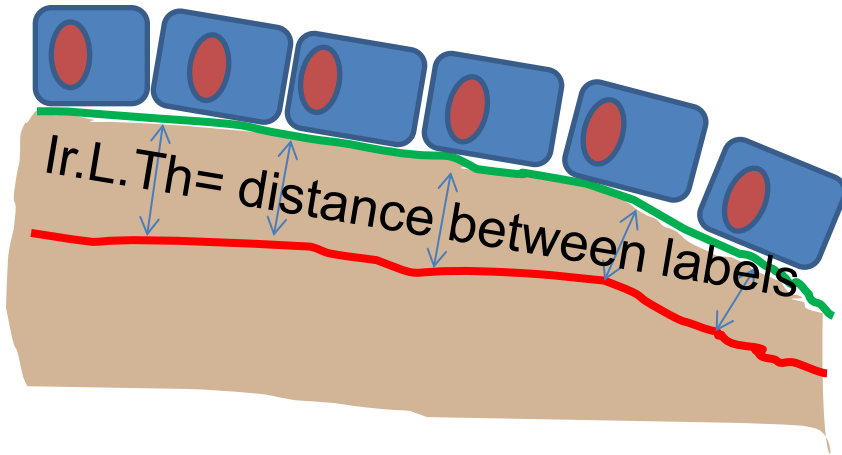
- Mineralizing surface $MS = (dLS + sLS/2)/BS$
 - the extent of bone surface actively mineralizing
- Mineral apposition rate $MAR = Ir.L.Th/Ir.L.t$
 - distance between the labels divided by time between labels
- Bone formation rate $BFR = MAR * (MS/BS)$
 - multiplies the MAR by the fraction of bone surface that is labeled

MAR vs BFR

$$\text{MAR} = \text{Ir.L.Th} / \text{Ir.L.t}$$

Ir.L.t = time between injection of labels

MAR is the rate at which OBs are making matrix, which calcifies at a constant rate and incorporates the labels. Thus, it measures the average activity of individual OBs in your section.



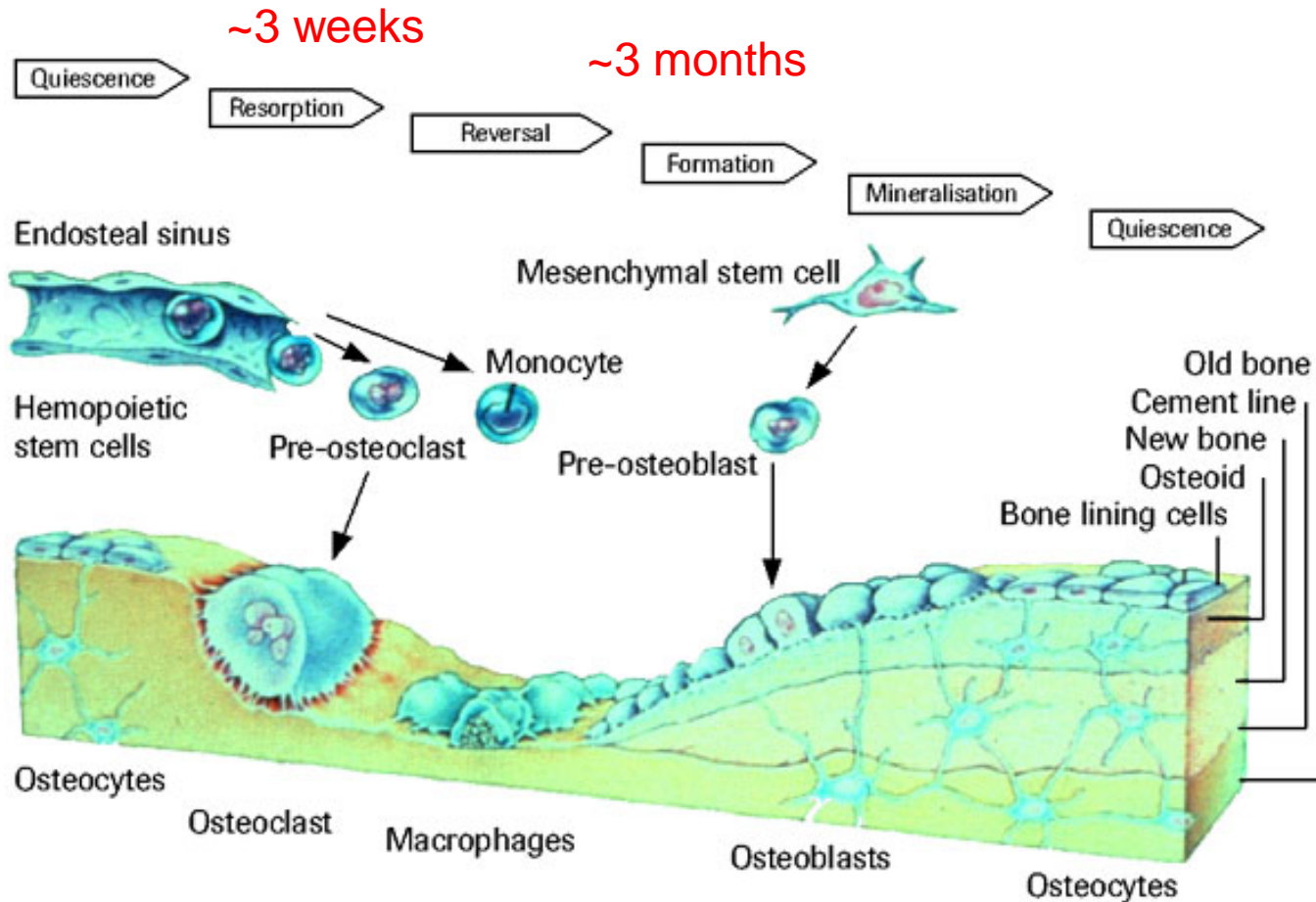
$$\text{BFR} = \text{MAR} * (\text{MS}/\text{BS})$$

BFR takes into account how much of the bone surface is actively mineralizing, which depends on the number of OBs that are active. It multiplies the average work of each OB by the fraction of bone surface with active OBs, giving a “team” output.

Bone formation rates

- There is a BFR for every possible referent (OS, BS, BV, TV) but usually we use BS
- ASBMR taskforce paper says
 - BFR/BS seems most logical when considering hormonal effects on bone remodeling
 - BFR/BV is equivalent to bone turnover rate, which determines bone age and various age-dependent properties of bone
 - BFR/TV seems most logical when considering biochemical markers of bone remodeling because the entire tissue is perfused and contributes its products to circulation

Bone Remodeling Cycle



For humans

Issues related to low bone turnover

- Recent studies of potent bisphosphonates and denosumab have run into issues where the dynamic labeling is very low, with no double labels with which to calculate BFR or MAR
- Need to state # samples with double labels, only single, or no labels, and the method of dealing with single labels
 - MAR can be recorded as missing data or a value assigned (imputed)
 - Either 0.1 $\mu\text{m}/\text{d}$ (lowest measured MAR) or 0.3 $\mu\text{m}/\text{d}$ (lowest average MAR)
 - How you include the samples with no double label has a big effect on the group averages

Effects of Denosumab on Bone Histomorphometry: The FREEDOM and STAND Studies

[J Bone Miner Res.](#) 2010 Oct;25(10):2256-65. doi: 10.1002/jbmr.149.

Ian R Reid,¹ Paul D Miller,² Jacques P Brown,³ David L Kendler,⁴ Astrid Fahrleitner-Pammer,⁵ Ivo Valter,⁶ Katre Maasalu,⁷ Michael A Bolognese,⁸ Grattan Woodson,⁹ Henry Bone,¹⁰ Beiyong Ding,¹¹ Rachel B Wagman,¹² Javier San Martin,¹¹ Michael S Ominsky,¹¹ and David W Dempster¹³ on behalf of the Denosumab Phase 3 Bone Histology Study Group

If the first unstained section did not show tetracycline fluorescence, then every tenth section was examined until tetracycline fluorescence was found. This section was used as the index slide, and the three groups of slides just described were prepared at 250- μm intervals. Examination at 50- μm intervals was continued throughout the entire biopsy, if necessary, to identify tetracycline labels (extended label search).

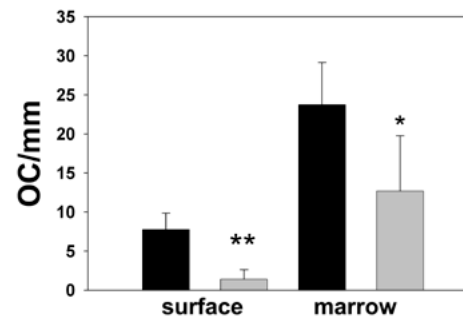
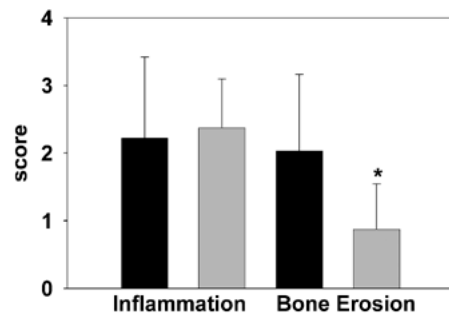
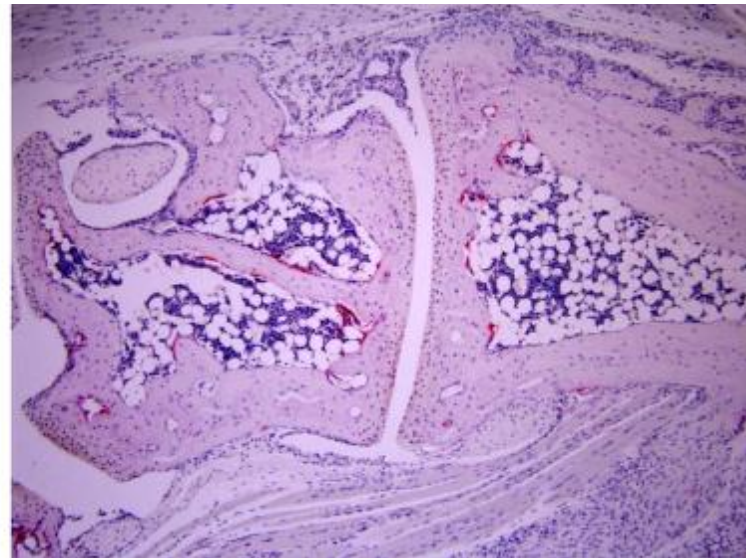
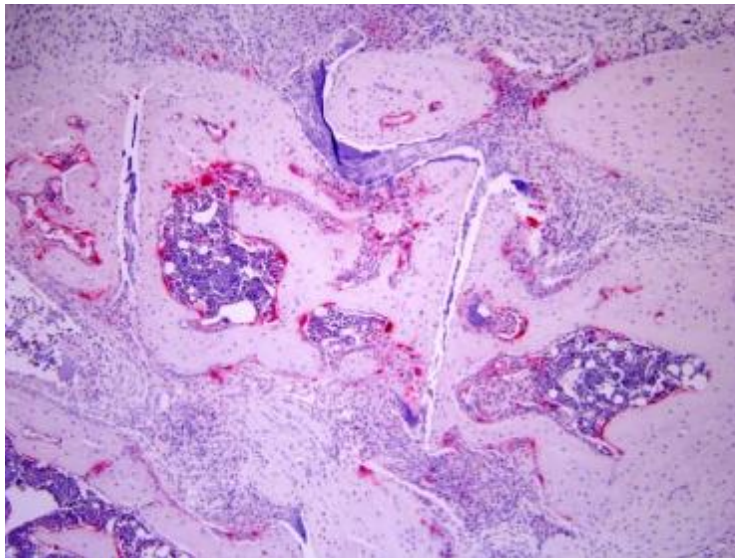
Analysis included data from all subjects who received one or more doses of investigational product and had one or more evaluable biopsies during the study. For biopsies with only single labels in trabecular bone within the measurement field of 20 mm², a value of 0.3 $\mu\text{m}/\text{day}$ was imputed for mineral apposition rate and used to derive other dynamic parameters, as recommended by Hauge and Foldes.^(18,19) Biopsies with single and/or double labels that were located only in cortical bone or were outside the measurement field in trabecular bone were excluded from these analyses. Between-group treatment differences for histomorphometric and μCT variables were evaluated using the Wilcoxon rank-sum test.

Examples of histomorphometry in disease models

Histomorphometry in Inflammatory Arthritis

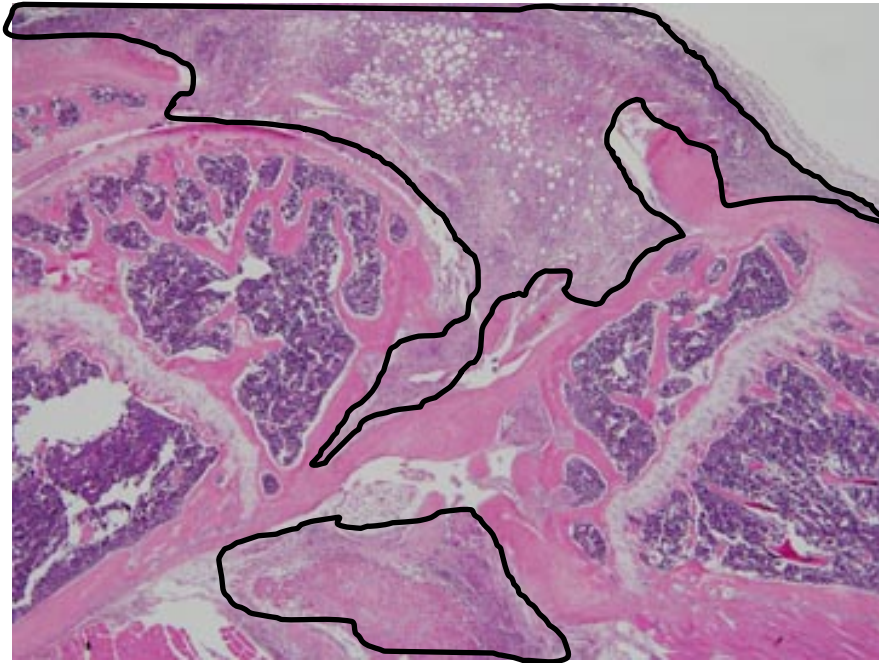
WT

NIK^{-/-}

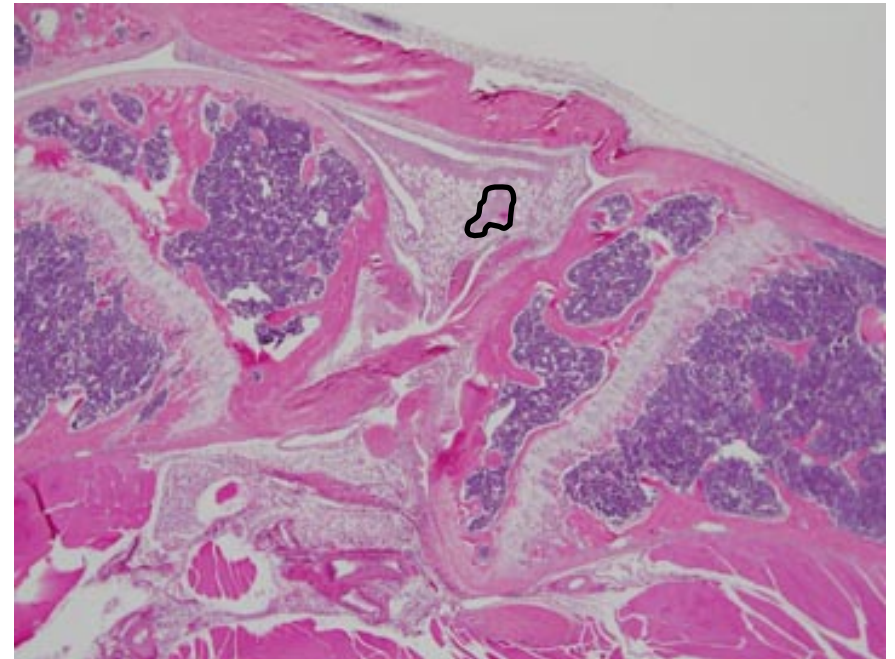


Antigen-induced arthritis

WT



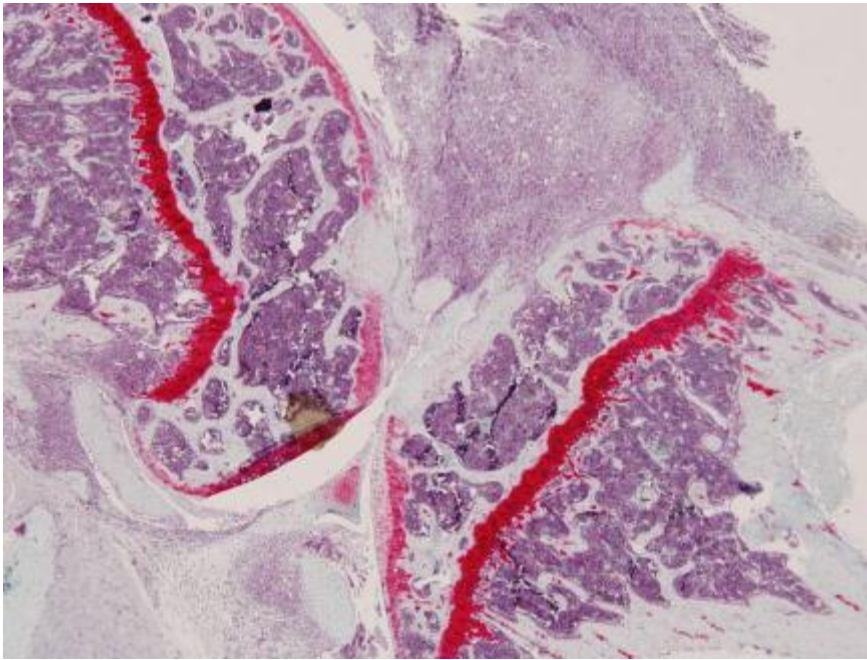
KO



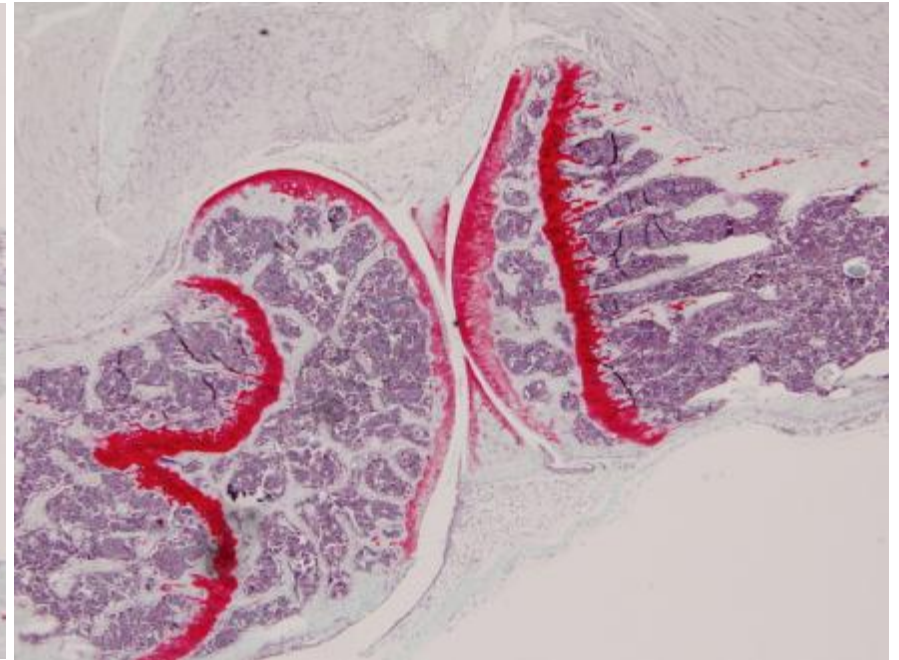
quantify area with inflammation

Antigen-induced arthritis

WT

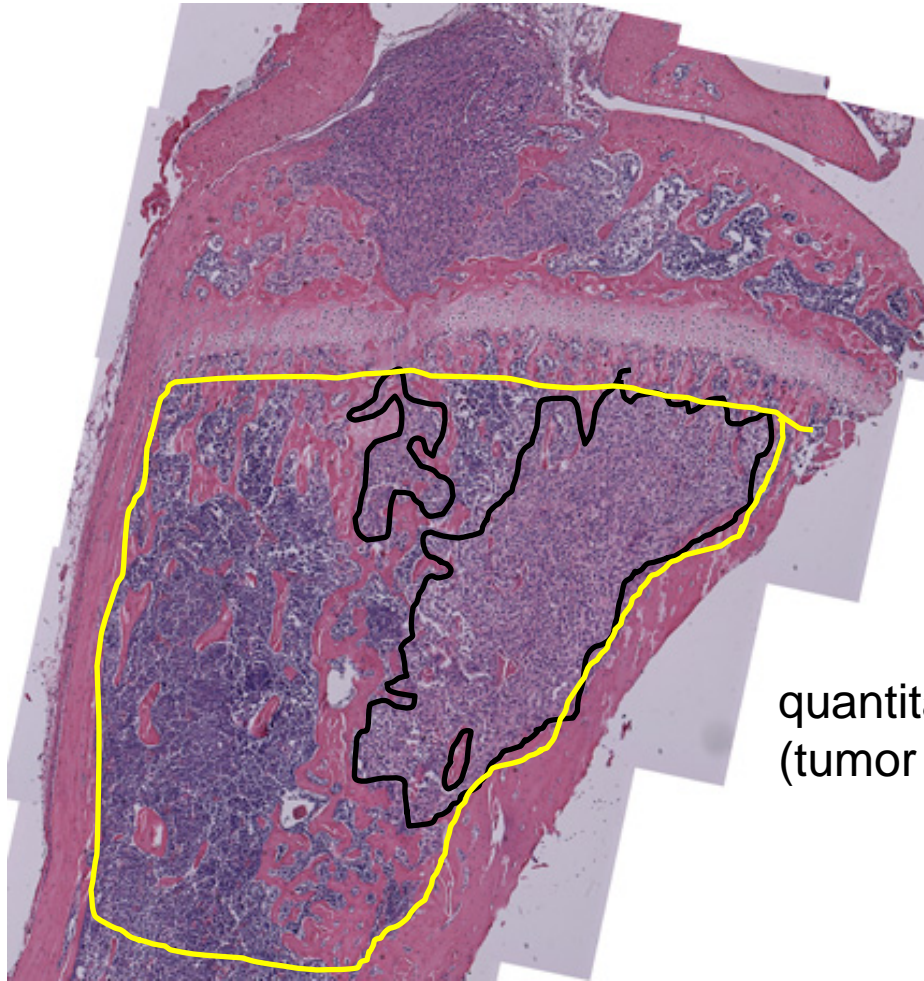


KO



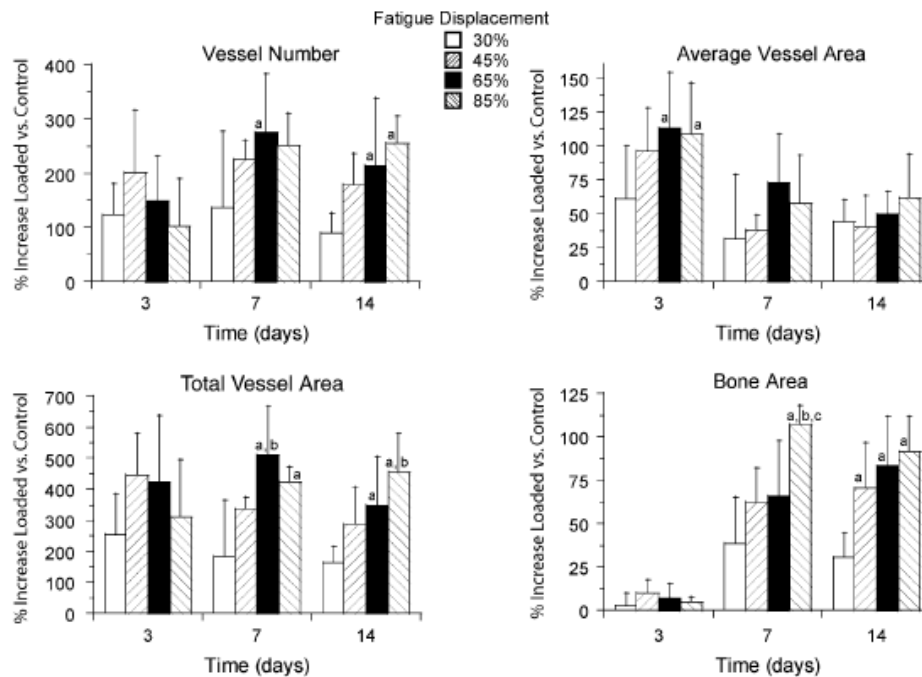
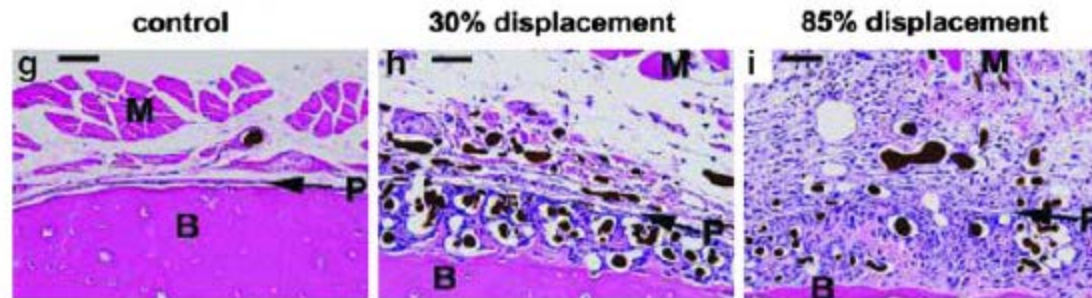
quantitate extent of safrO staining on articular surface

Bone metastasis



quantitate tumor area/tissue area
(tumor volume/tissue volume)

Analysis of fracture model looking at vessels



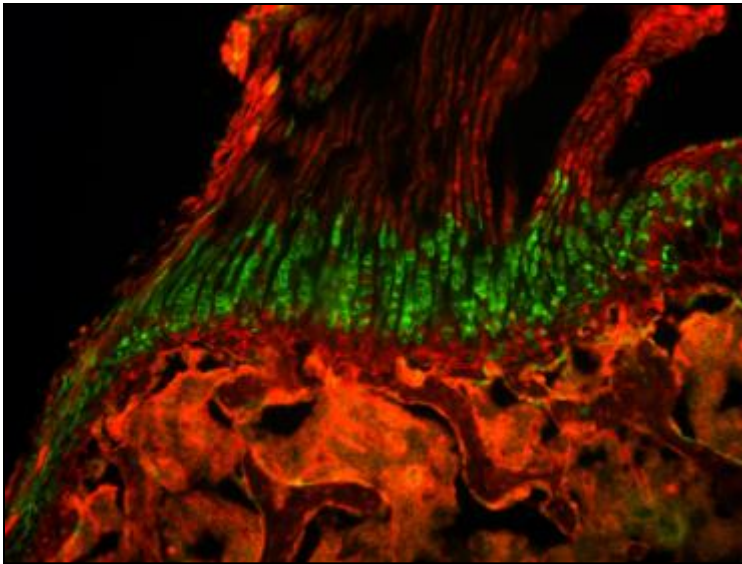
Silva group

Newer options – frozen sections

- Cryojane tape transfer system (Leica)
 - Tape is applied to tissue block and knife cuts behind the tape, leaving your section adherent to the tape
 - Tape is rolled onto a special adhesive slide and tissue is transferred to slide with a zap (or several) of UV
 - Nondecalcified tissue does not transfer well, so this is best for (at least partially) decalcified samples
 - Can be used for histo stains, IHC, or IF
- Cryofilm tape (Section labs)
 - Film is applied to block and knife cuts behind
 - Film is then glued to a regular slide, tissue side up
 - Tissue can be stained, then coverslipped
 - Special LCM film is available for laser capture, with special slides (with a hole in the middle)

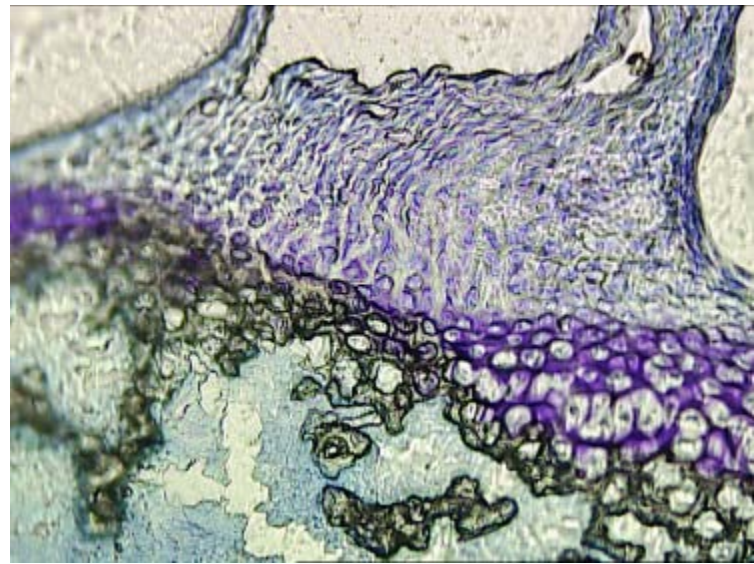
Cryojane/cryofilm

Cryojane



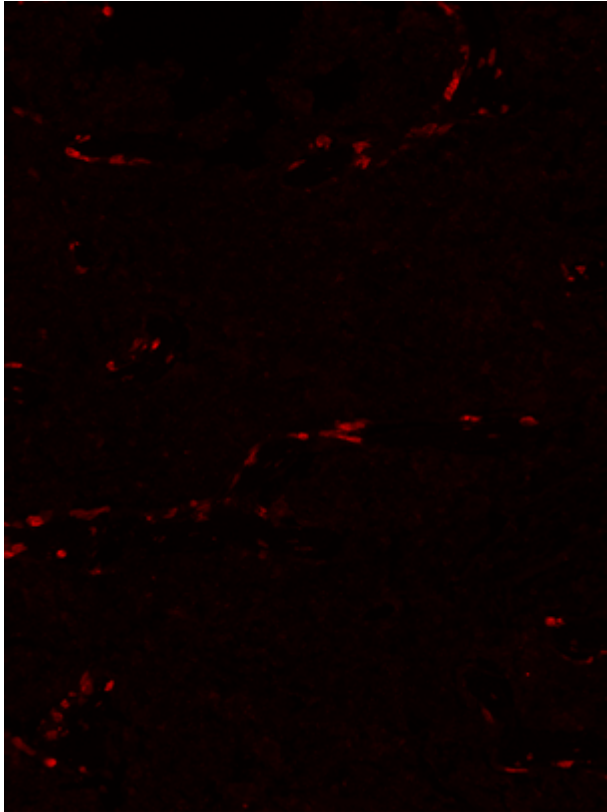
Frozen section of fixed, decalcified and frozen tissue observed without further staining under fluorescence, mTmG reporter

Cryofilm

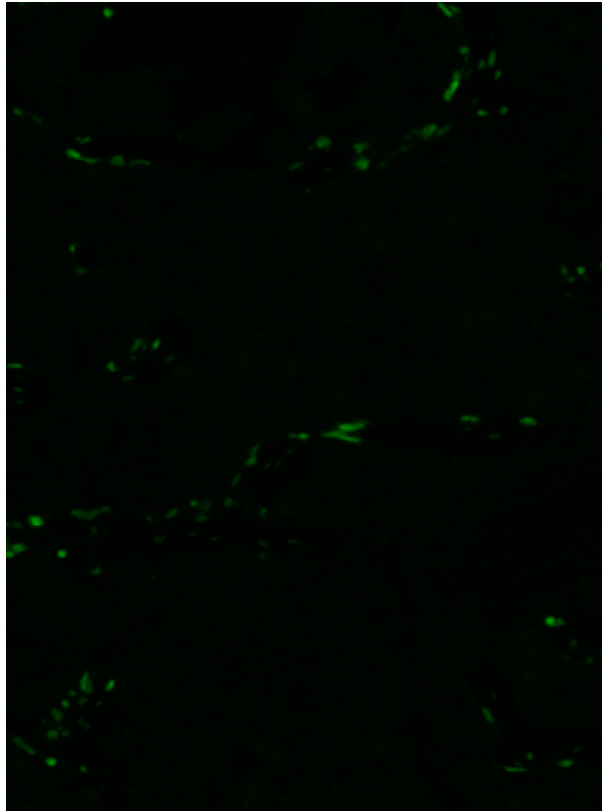


Unfixed, nondecalcified frozen section, toluidine stain

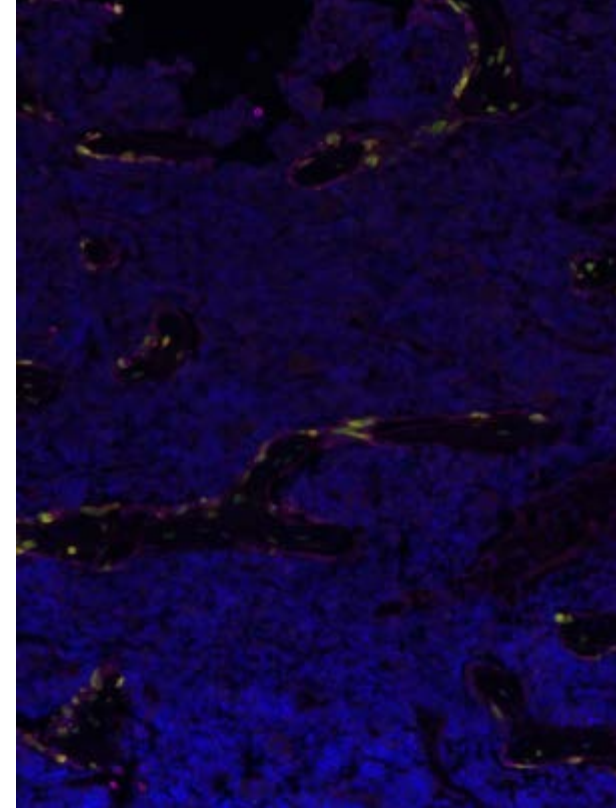
Cryofilm example



IF, osteocalcin

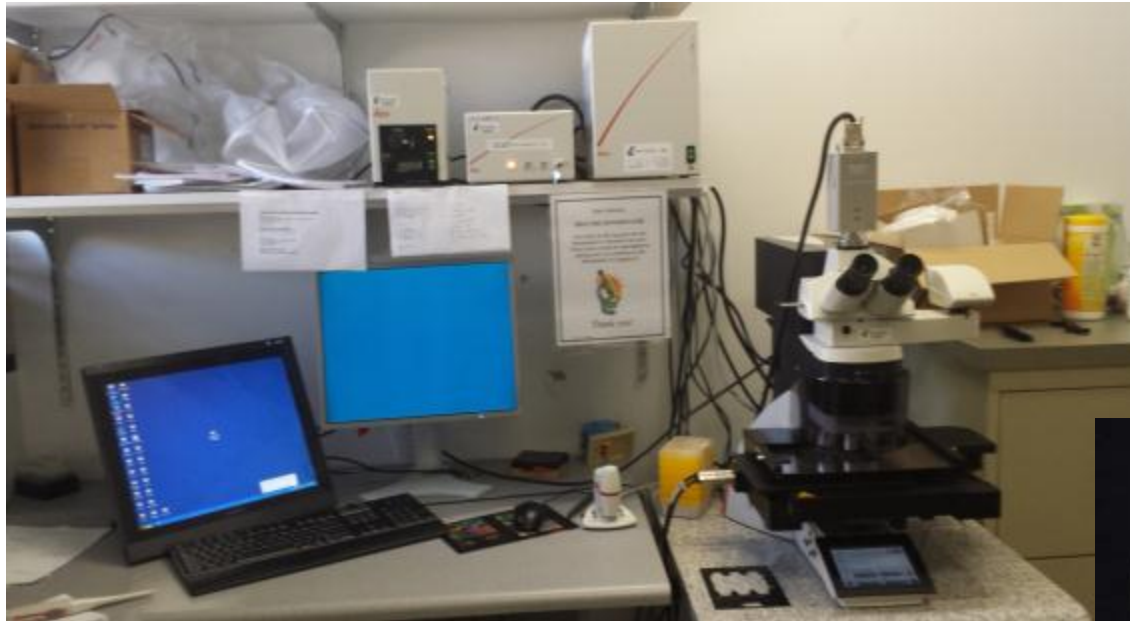


GFP transgene, no stain

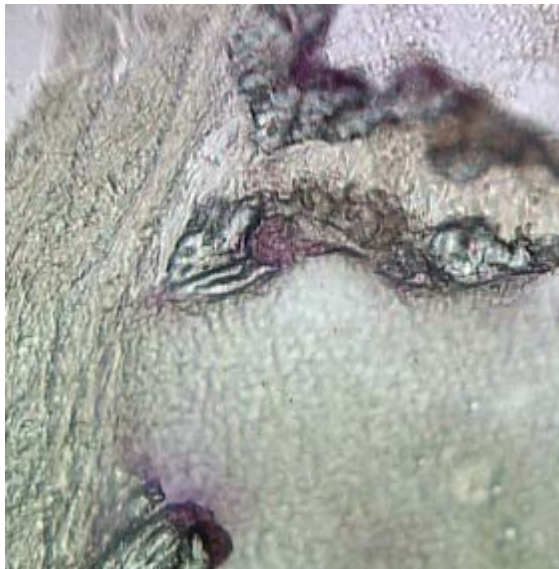


Overlay, with Hoechst

Laser Capture Microdissection



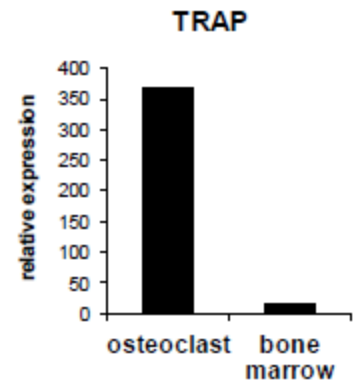
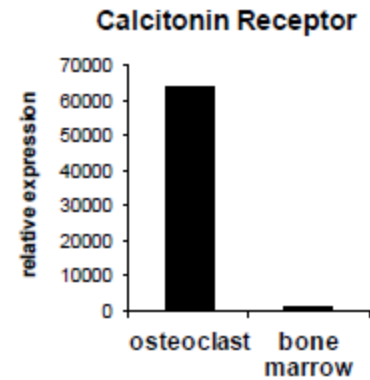
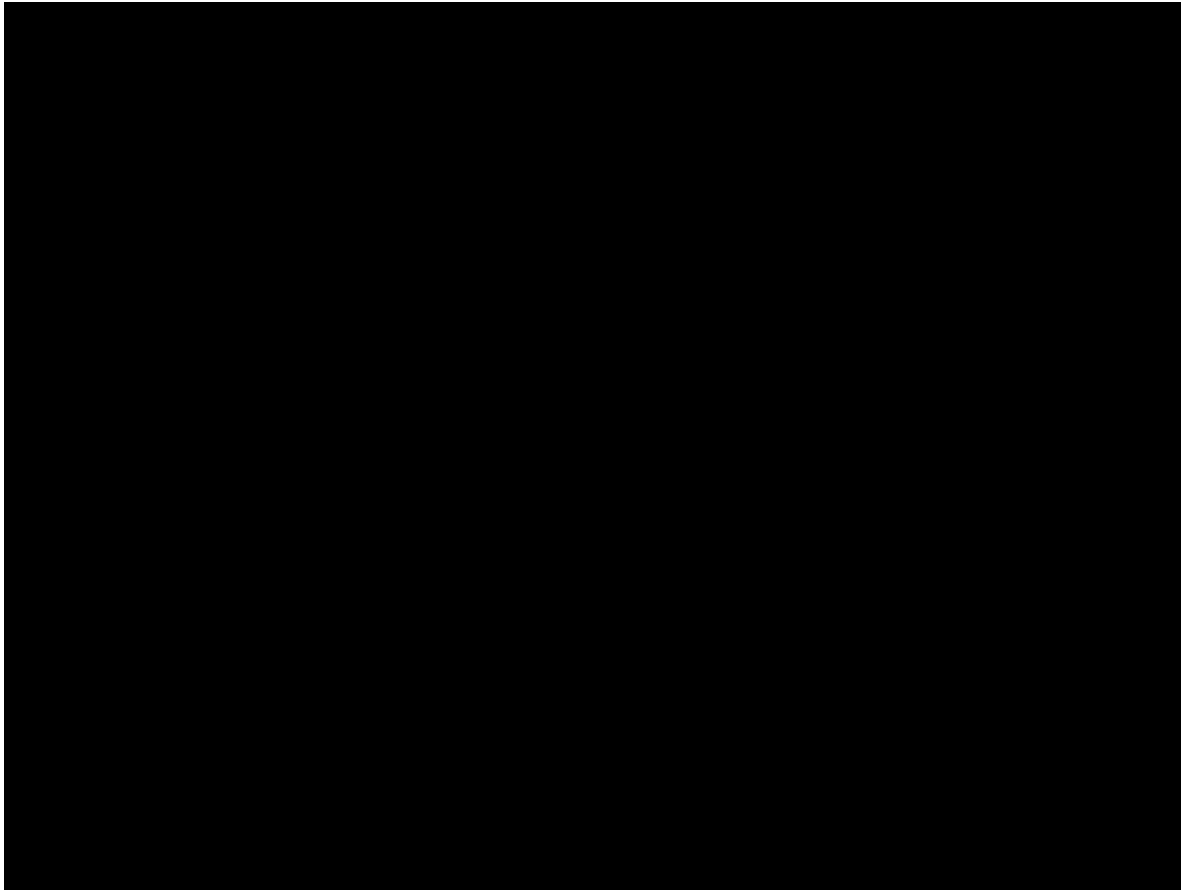
Laser beam cuts tissue
and it falls into cap



Brief TRAP stain



Draw a contour, and cut with laser



RNA suitable for qPCR
Testing for RNAseq now

From Chang Yang

Useful references

Standardized Nomenclature, Symbols, and Units for Bone Histomorphometry: A 2012 Update of the Report of the ASBMR Histomorphometry Nomenclature Committee

David W Dempster,^{1,2} Juliet E Compston,³ Marc K Drezner,⁴ Francis H Glorieux,⁵ John A Kanis,⁶ Hartmut Malluche,⁷ Pierre J Meunier,⁸ Susan M Ott,⁹ Robert R Recker,¹⁰ and A Michael Parfitt¹¹

J Bone Miner Res. 2013 January ; 28(1): 2–17. doi:10.1002/jbmr.1805.

Histomorphometry in Rodents

Reinhold G. Erben and Martin Glösmann

Miep H. Helfrich and Stuart H. Ralston (eds.), *Bone Research Protocols*, Methods in Molecular Biology, vol. 816, DOI 10.1007/978-1-61779-415-5_19, © Springer Science+Business Media, LLC 2012

Thank You !!

Less commonly used terms

- More relevant in clinical studies

Adjusted apposition rate

- $Aj.AR = MAR * MS / OS$
 - Represents the mineral apposition rate averaged over the osteoid surface
- In a steady state and in the absence of osteomalacia (mineralization defect), Aj.AR is the best estimate of the mean rate of osteoid apposition, and should be the same as the rate of mineral apposition
- The mineralization lag time (how long it takes to mineralize a newly deposited hunk of osteoid) is the osteoid thickness divided by the Aj.AR
 - $Mlt = O.Th / Aj.AR$
 - Osteomalacia has been defined as $Mlt > 100 \text{ d}$

Wall width (W.Wi) and Formation period (FP)

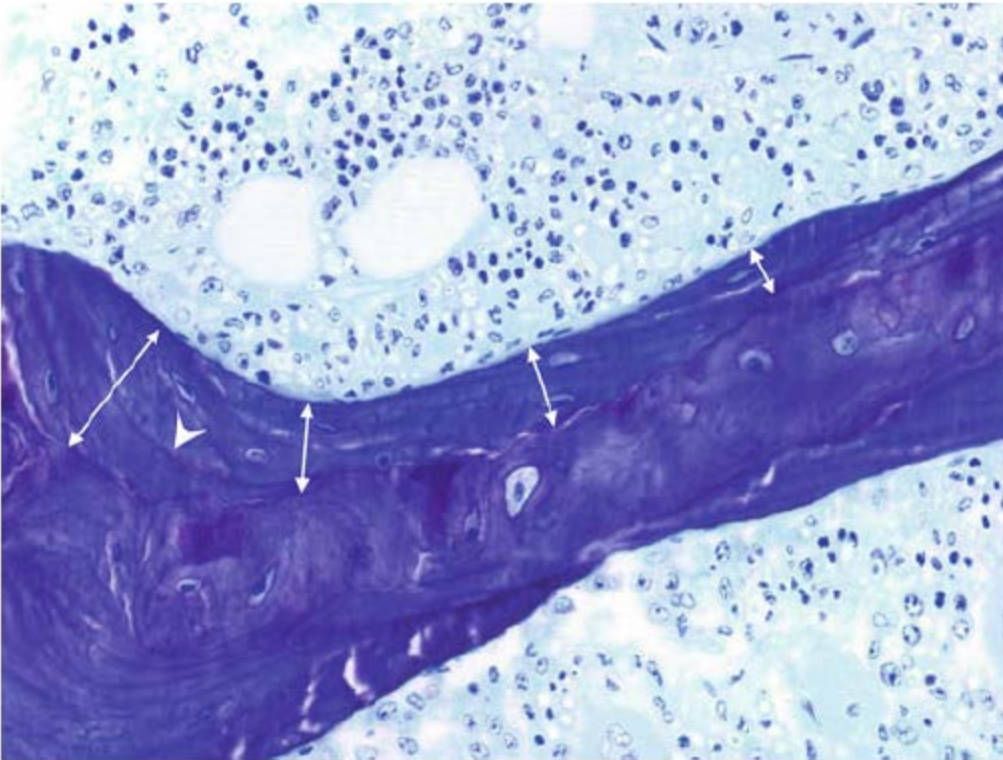


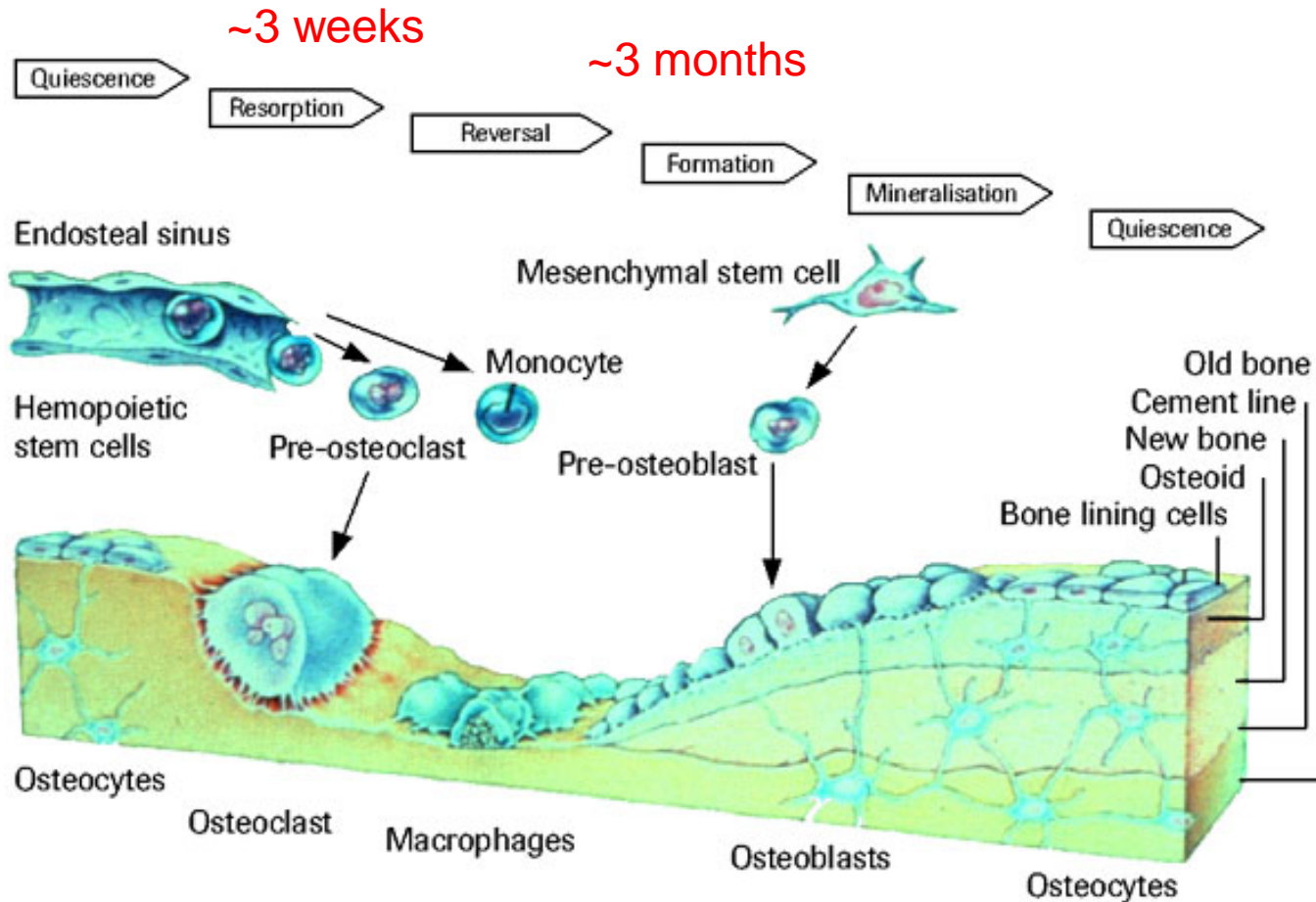
Fig. 7. Measurement of wall width for remodeling-based histomorphometric parameters. Wall width is defined as the mean distance between the bone surface and the scalloped reversal line in individual completed remodeling units. A smooth arrest line, indicating a temporary stop in osteoblastic bone formation within a remodeling unit, is marked by an arrowhead. Five-micron-thick section of a rat lumbar vertebra surface-stained with cement line stain. Original magnification $\times 400$.

- Wall width is used to determine the amount of bone added during a remodeling cycle
- The idea is that the surface extent of a certain activity is proportional to the time occupied by the activity
- Measure at least 15 units/sample
- Convert to thickness (W.Th in 3-D) by multiplying by $\pi/4$
- $FP = W.Th / MAR$ in rodents or $W.Th / A_j.AR$ for humans

Activation frequency (Ac.f)

- The Ac.f is the probability that a new cycle of remodeling will occur at any point on the surface by the event of activation
 - It is defined as $1/Tt.P$ (total period) or the reciprocal of the total time it takes for a remodeling cycle
 - $Tt.P = FP*(BS/OS)$
 - We don't have a direct way of measuring the resorption or reversal periods, although there are formulas based on Oc.S and ES

Bone Remodeling Cycle



For humans