SOLID Heritage Bone Diagenesis

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Image Reznikov, N., Bilton, M., Lari, L., Stevens, M. M., & Kröger, R. (2018). Fractal-like hierarchical organization of bone begins at the nanoscale. *Science*, **360**(6388). https://doi.org/10.1126/science.aao2189



Taphonomy

Skeletal tissues may be either exposed on the ground surface (Biostratinomy) or buried in various burial environments (Diagenesis)

- Biostratinomy covers the pre-burial histories of bones.
- Diagenesis covers the post-burial histories of bones.





What is Diagenesis?

- Post-mortem each object seeks to reach equilibrium with the conditions of its burial environment.
- A series of physical, chemical, and biological processes affect its preservation state. These are called diagenetic trajectories or diagenetic pathways.
- Three diagenetic pathways are identified for bone:
 - 1. Biological deterioration of the composite (i.e. microbial attack).
 - 2. Chemical deterioration of the inorganic component (i.e. dissolution).
 - 3. Chemical deterioration of the organic component (i.e. hydrolysis).

Diagenetic Parameters

State of preservation of bone characterized by 12 simple measurements

• Microstructure

Histology: Oxford Histological Index; Collagen Birefringence Index; Cracking; Inclusions

• Physical

Porosity: Mercury Porosimetry

Biomechanical properties: density measurements (micro- and nano-indentation)

• Inorganic

IRSF (crystallinity); C:P (Carbonate-to-Phosphate); Identification of other mineral phases

• Organic

%N of Whole bone; % 'Collagen'; C:N ratio of 'Collagen'

Histology



Wedl tunnelling

Is this just expanded and stained canaliculi?

Budded



Histology: Quantitative Assessment

Oxford Histological Index				
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	OXIDIU	1 115101	USICUI	ITIGEX

0 <5 No original features identifiable, except that 0 Haversian canals may be present 0.5	ensity of Birefringence
0.5	Absent
	Reduced
Haversian canals present, small areas of1Normal1<15	, comparable to fresh bone
is preserved by the pattern of destructive foci Jans et a	I. (2002), Archaeometry 44 (3):343–352
2 <50 Some lamellate structure is preserved between the destructive foci	
3 >50 Some osteocyte lacunae preserved	
4 >85 Bone is fairly well preserved, with minor amounts of destructive foci	
5 >95 Very well preserved, virtually indistinguishable from modern bone	

Birefringence Index

Millard (2001), The deterioration of bone. In *Handbook of Archaeological Sciences*, D. R. Brothwell, A. M. Pollard (eds.). Wiley: New York. 637–674.

Histology: Qualitative Assessment



- Fungi (and cyanobacteria in marine environments) create branched tunnels in the bone matrix called Wedl tunnels.
- Bacteria reorganize the mineral content of bone by creating microscopic focal destructions (MFD) called non-Wedl tunnels.



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Figure 1.2 (a) Transmitted light image of medieval human bone from Trondheim, Norway. Histological preservation is excellent, but staining around the central osteon illustrates the fine canalicular network that connects the tissues with the soil environment. (b) The section viewed in polarized light with a quarter-lambda plate. The spectacular birefringence arises from the alignment of collagen fibrils and HAP in the bone lamellae

Histology

Cracking

- Micro-fissures due to deterioration of the inorganic and/or organic components.
- Decoupling of mineral-collagen bond, demineralization or thermal stress.
- Large fissures maybe due to physical stress.



Jans (2005), Histological Characterisation of Diagenetic Alteration of Archaeological Bone. PhD Thesis, Vrije Universiteit: Amsterdam

Histology

Inclusions - Pyrite framboids







Diagenetic Parameters

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%N of Whole bone; % 'Collagen'; C:N ratio of 'Collagen'

Porosity

Mercury intrusion porosimetry: to measure intrusion of mercury into a porous structure under stringently controlled pressures (pore size distribution, volume, diameter, etc.).



Porosity changes modern cow bone deproteinated cow bone deproteinated bird bone archaeological bone



Porosity: pH > 6.0

increase> 0.2 µm



Dissolution removes mineral

- Why no organic matter?
- Removal of mineral exposes collagen



Fossilization: Apigliano (& Çatalhöyük?)



Patterns of microbial attack





Porosimetry trace





Image analysis: Gordon Turner-Walker





Human vs. animal bone



Miranda Jans

Churchyards







Animal vs. human bone

- Human bone
 - Predominately microbial attack
 - Burial with intact blood supply
 - Putrefaction of intestine
 - Migration of gut bacteria *Clostrida* sp. into blood then bone

The aller

- Animal bone
 - Fungal attack (Wedl)
 - No putrefaction
 - Processed bone
 - [100% of interred animals (9) have microbial attack]

Porosity: well preserved bone



Mycorrhizal fungi



Boiling bone?



Ç

Burnt bone?



Principal Components Analysis



axis 2 (22%)

PCA



Bone corrosion





Bronze Age Britain?



Booth, T. J., Chamberlain, A. T., & Pearson, M. P. (2015). Mummification in Bronze Age Britain. Antiquity, 89(347), 1155–1173.

Leads to generalised ideas...

Oxford Histological Index

Index	Approximate % of intact bone	Description
0	<5	No original features identifiable, except that Haversian canals may be present
1	<15	Haversian canals present, small areas of well-preserved bone present, or lamellate structure is preserved by the pattern of destructive foci
2	<50	Some lamellate structure is preserved between the destructive foci
3	>50	Some osteocyte lacunae preserved
4	>85	Bone is fairly well preserved, with minor amounts of destructive foci
5	>95	Very well preserved, virtually indistinguishable from modern bone



Poor histology



1

0



Well preserved histology

What can you see here?

Computer Tomography (CT)



Image Rich Magdwich Cardiff



Image Rich Magdwich Cardiff

CT imaging: Infant bone

Half of the infant samples studied here were free from bacterial bioerosion, further suggesting that histological analysis can be used to identify archaeological remains of stillborn and short-lived infants.

The samples that were not tunnelled had "bright inclusions" in them. Probably pyrite framboids. Thus, probably in anoxic conditions.





Experimental burials

Anne-Marie Høier Eriksen,

Experimental

- (a) Raw bone fragments
- (b) cutting fragments
- (c) spears with raw,boiled and baked bone fragments were embedded at the two submerged study environments; half the fragments were embedded into the sediment and the other half were suspended in the water column
- (d) terrestrial environment the fragments were placed at the bottom of a 30 cm-deep trench;
- (e) sea-bed spears overgrown with algae after 28 weeks
- (f) bones at the tidal zone were left on the shore and subjected to wet-dry cycles.



Eriksen, A. M. H., Matthiesen, H., Kontopoulos, I., Gregory, D., Snoeck, C., Zhang, G., Collins, M. J., & Gilbert, M. T. P. (2020). Rapid loss of endogenous DNA in pig bone buried in five different environments. Archaeometry, **62**(4), 827–846.

Submerged gyttja

Baked bone, Submerged gyttja, 52 weeks.

- chemical demineralisation on the periosteal surface.
- cyanobacterial attack.
- enlarged and ragged osteocyte lacunae and enlarged canaliculi.



Tidal Zone: Cyanobacteria



Tidal zone, buried for 52 weeks.

Enlarged area showing extensive cyanobacterial attack on the periosteal side. All samples at the tidal zone at 52 weeks showed this kind of damage.

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Terrestrial

Terrestrial sand, 52 weeks. Microbial tunnelling is observed from the periosteal border and to 200 µm depth.



DNA analysis of the microbial community

Eriksen, A. M. H., Nielsen, T. K., Matthiesen, H., Carøe, C., Hansen, L. H., Gregory, D. J., Turner-Walker, G., Collins, M. J., & Gilbert, M. T. P. (2020). Bone biodeterioration-The effect of marine and terrestrial depositional environments on early diagenesis and bone bacterial community. *PloS One*, **15**(10), e0240512.

Actinobacteria: Streptosporangium

Clostridiacea and Fusobacteriaceae in the two submerged sites,

both of which are known to be have members linked to collagenolytic enzymatic activity

Xanthomonadaceae (a Proteobacteria) found in the Terrestrial site,

has previously been found in degrading bone from individuals were placed outside on the ground surface to decompose naturally at the Anthropology Research Facility (ARF) at the University of Tennessee, Knoxville

Most common organism Actinobacteria Streptosporangium

Neanderthal bone

3.95 Gb of Neanderthal DNA isolated from the Vindija Neanderthal Vi33.16 fossil showed that 90% of about 50,000 rRNA gene sequence reads were of bacterial origin, of which Actinobacteria accounted for more than 75%.

Actinobacteria also represented more than 80% of the PCR-amplified 16S rRNA gene sequences from a cave sediment sample taken from the same G layer as the Neanderthal bone.

The bacterial DNA showed no signs of damage, and we hypothesize that it was derived from bacteria that have been enriched inside the bone.

Phylogeny of microbial collagenases.

Collagenase consensus sequences are coloured in red The actinobacterial clade is highlighted in yellow

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Collagenase

Dissolution removes mineral

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- Removal of mineral exposes collagen

SOLID

Henning and Scott

