

Paleogenetic analysis for the valorization of Cultural Heritage

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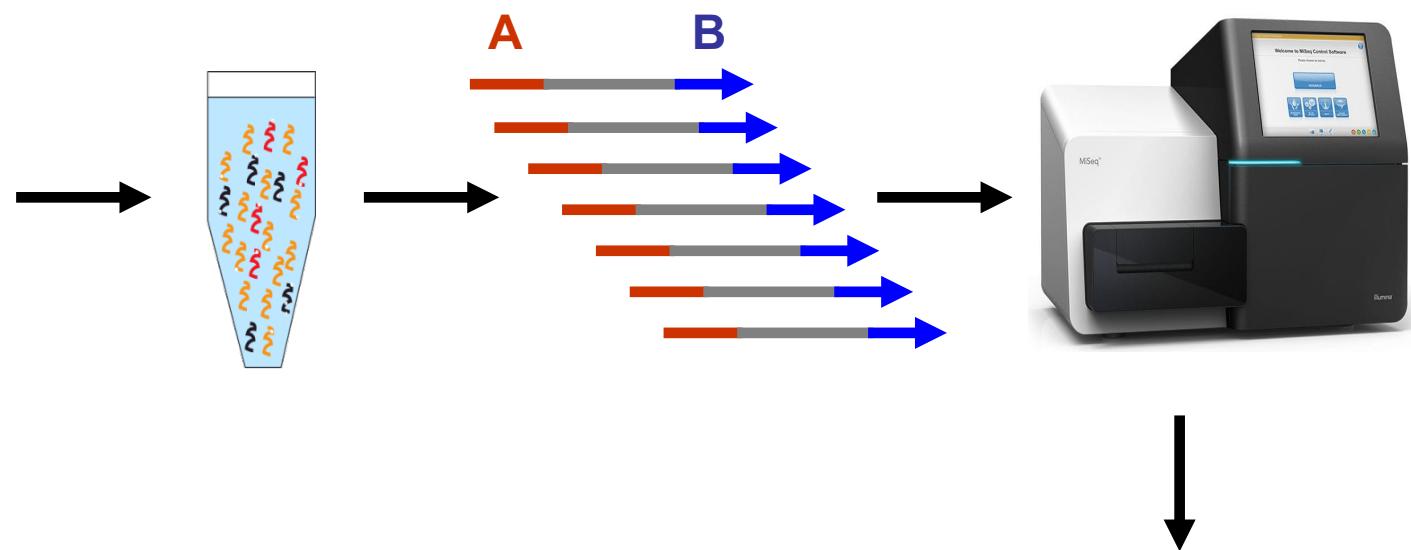
Laboratorio di Antropologia Molecolare e Paleogenetica



J.M Cecchi., R. Stanyon, *Il Museo di Storia Naturale dell'Università degli Studi di Firenze. Le collezioni antropologiche ed etnologiche*, Firenze University Press, 2014.



DNA analysis on ancient biological remains



- determination of species of origin
- individual identification of famous personalities
- phylogenetic relationships
- reconstructing life style and cultural contexts



Paleogenetic analysis



Human skeletal remains

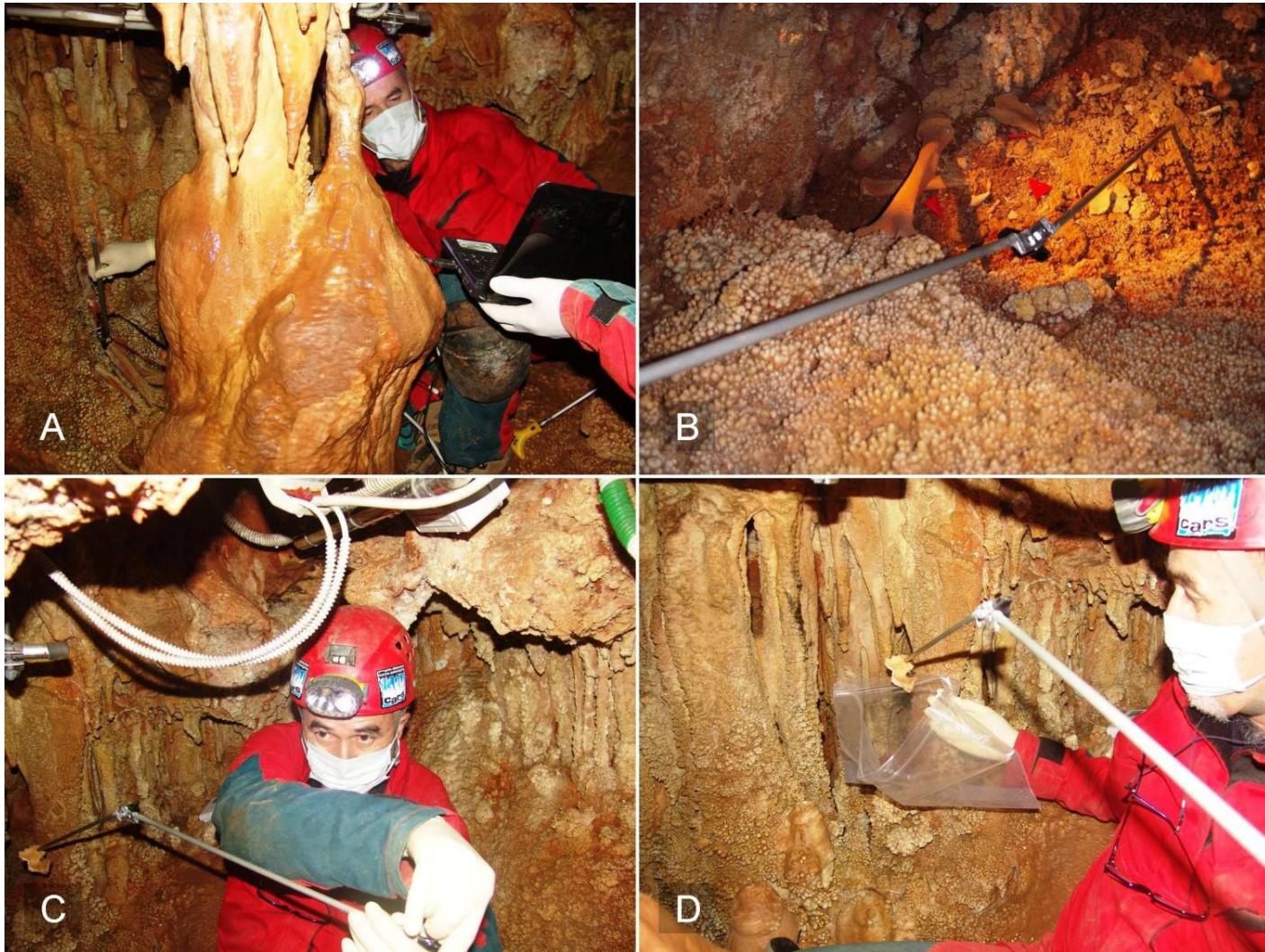




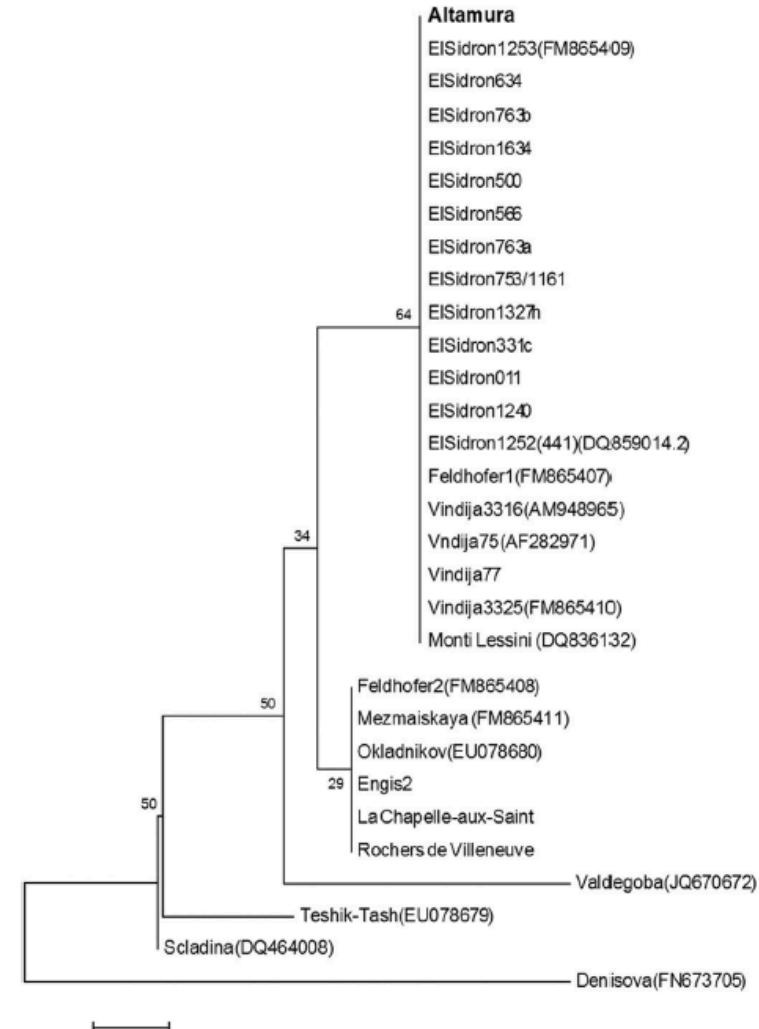
The human skeleton from Lamalunga, Altamura

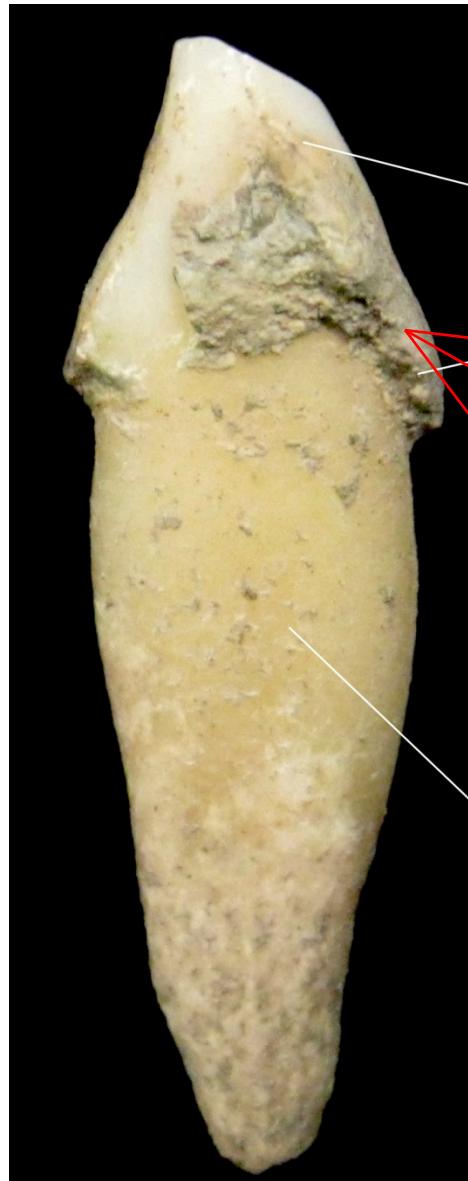


Removing a skeletal fragment from the cave

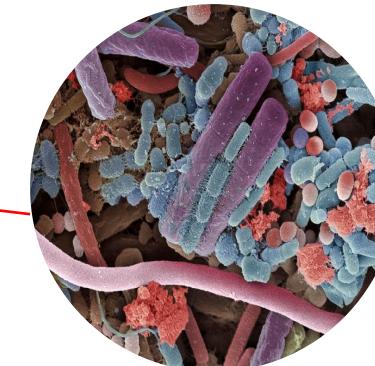


Results of the paleogenetic analysis





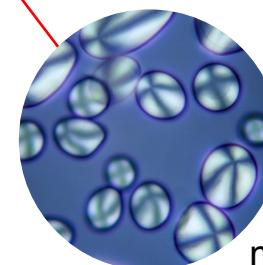
Ancient *dental calculus*



Bacterial flora



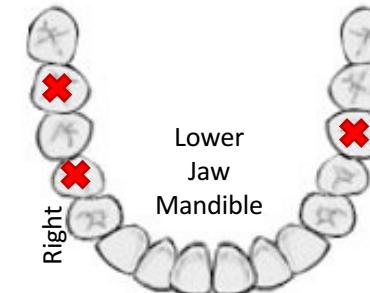
Human DNA from mucosa cells



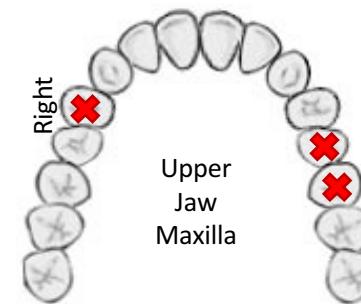
Plant microresidues

Porticus Octaviae (Rome), IX-XII sec. CE

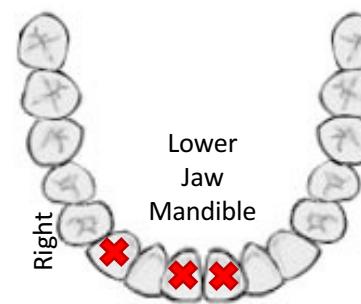
PO. 108-7



PO. Z. US44

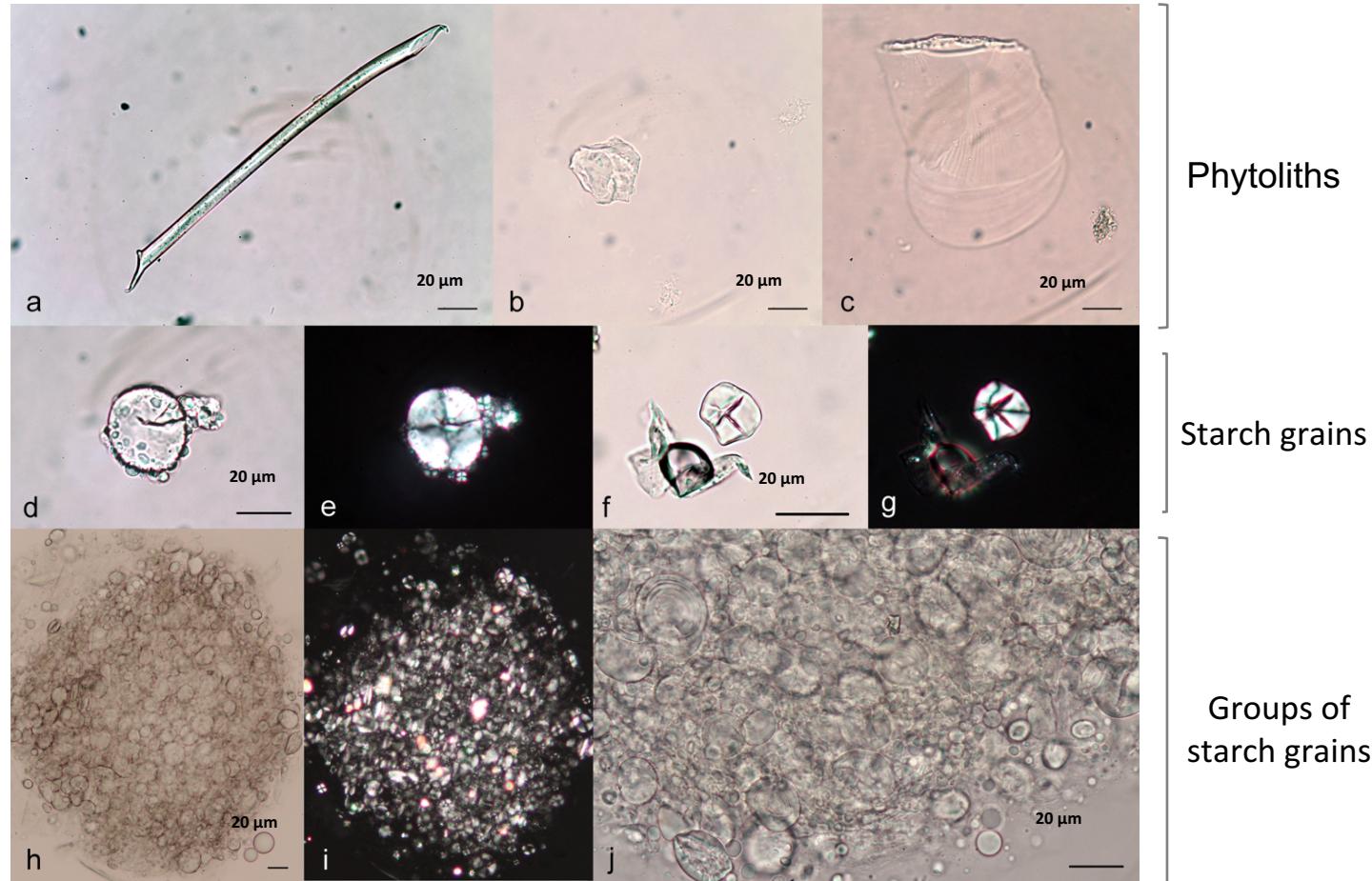


PO. 898-11

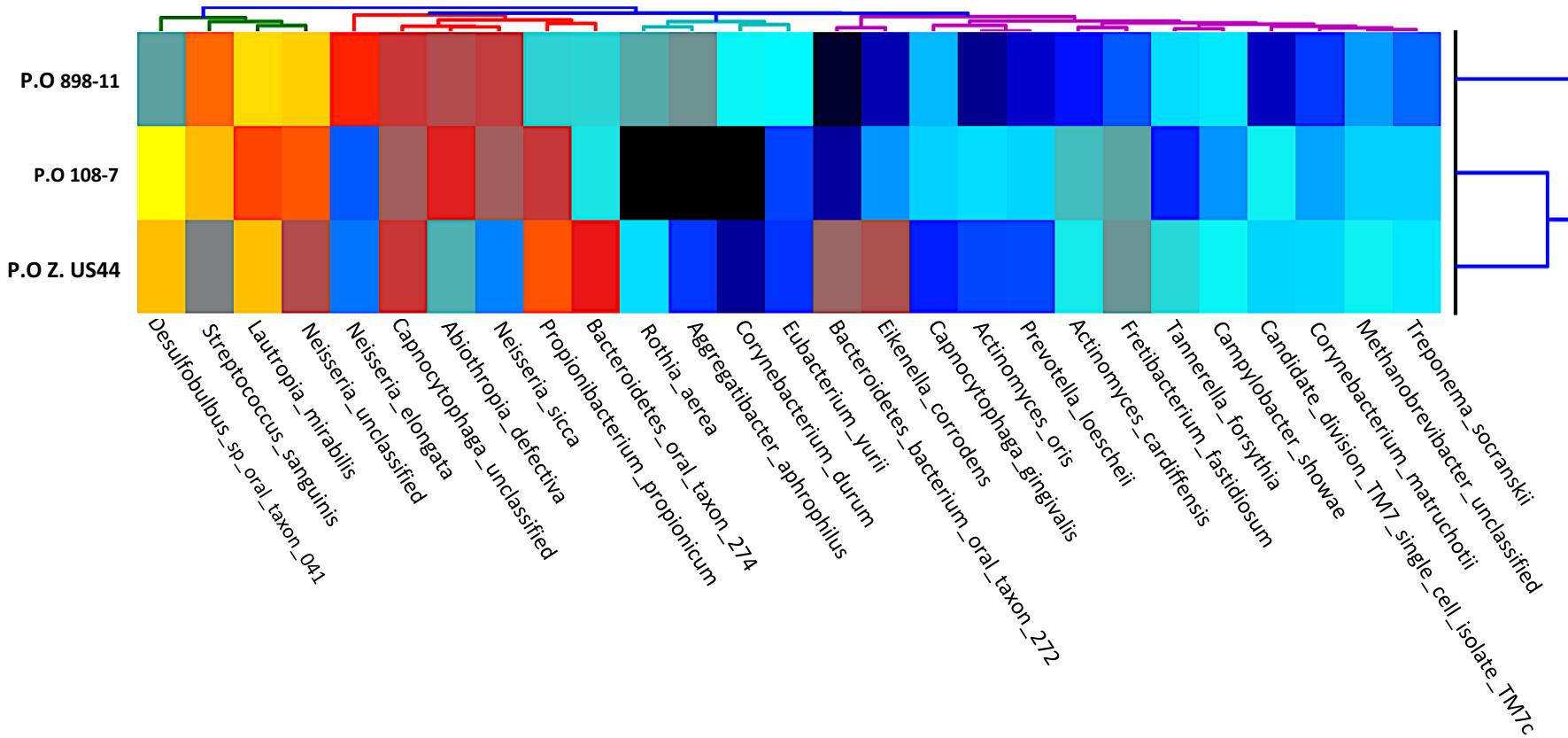


Dietary plant microresidues

- Poaceae, Panicoideae, Hordeum, Triticum

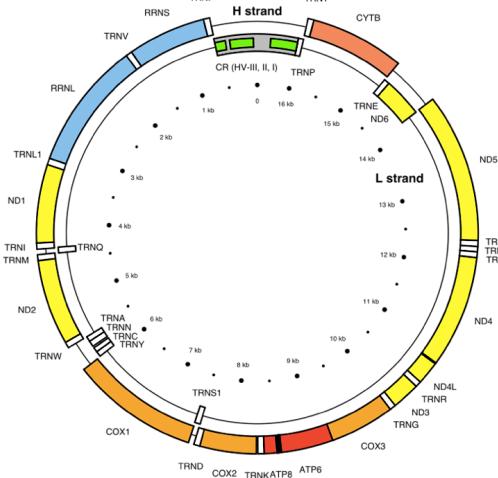


Oral microbiome composition

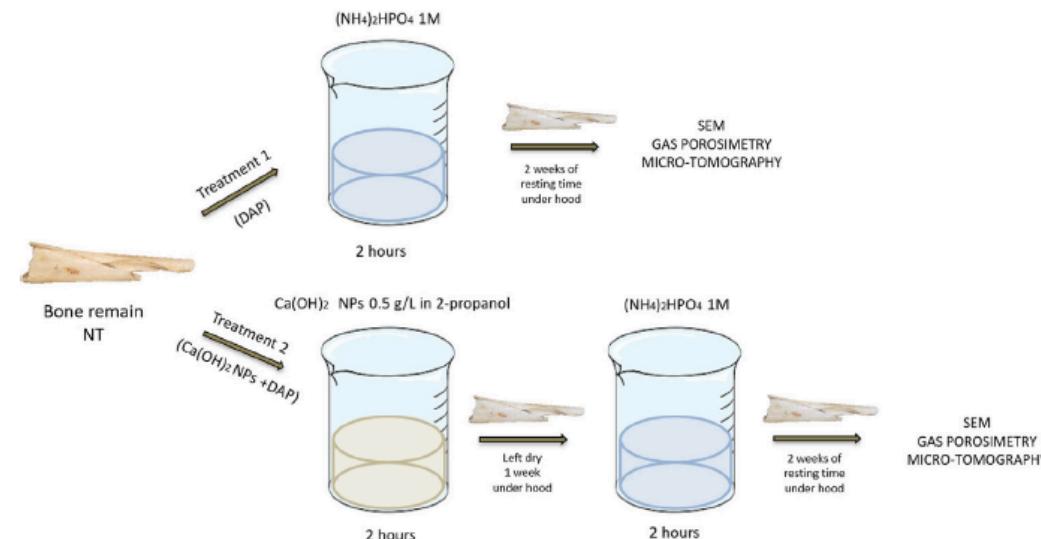


Complete human mitochondrial genomes

Sample ID	Hg	Total Variants	Variants	Missing positions
P.O 108-7	U5a1c2a1	26	A73G, A183G, A263G, A750G, A1438G, A2706G, A4769G, C7028T, A8860G, G9477A, C10544T, A11467G, G11719A, A12280G, A12308G, G12372A, T13617C, C14766T, A14793G, A15326G, C16192T, C16256T, C16270T, C16286T, C16320T, A16399G	15
P.O Z. US44	X2 + 225 + @16223	21	A73G, A153G, T195C, G225A, A263G, A750G, A1438G, G1719A, A2706G, A4769G, T6221C, C7028T, A8860G, G11719A, C12705T, A13966G, T14470C, T15090C, A15326G, C16278T, T16519C	10

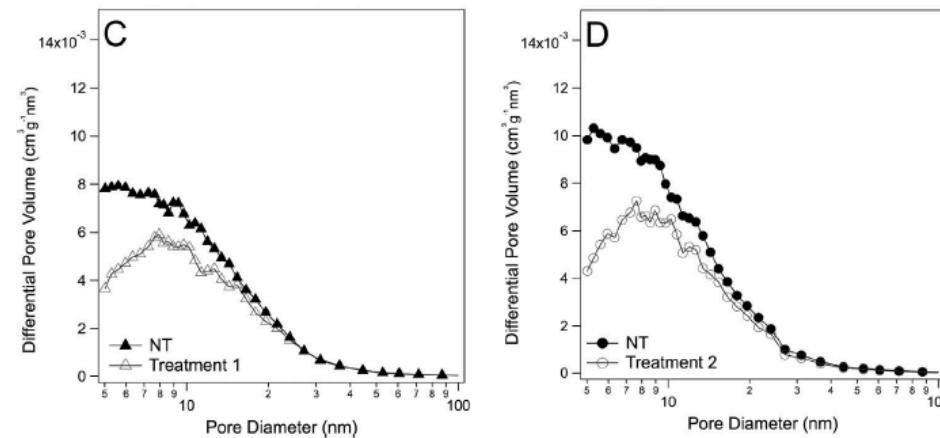


Restoration of ancient skeletal remains

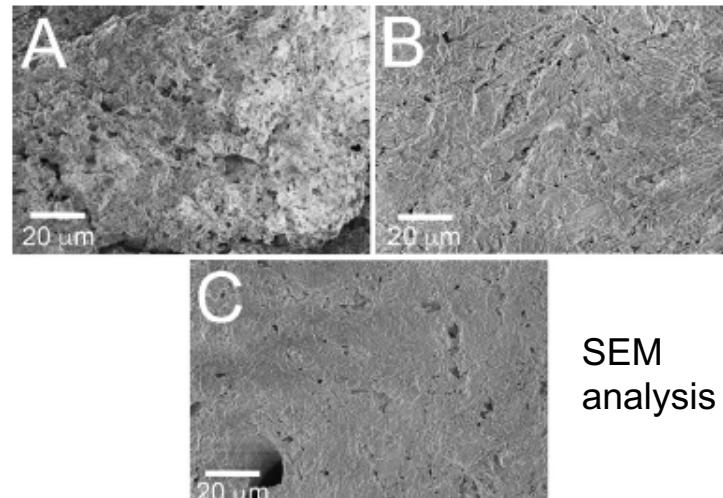
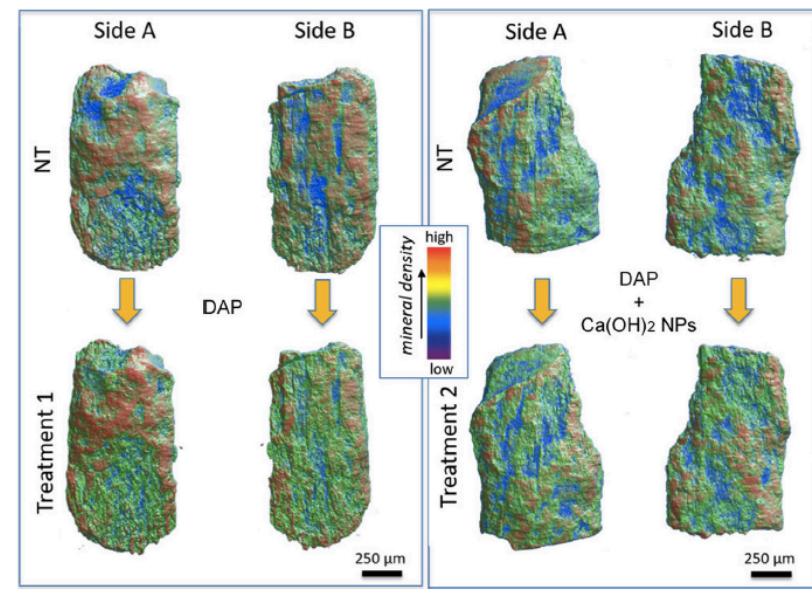


Scheme 1. Representative sketch describing the consolidation treatments on the bone remains. Consolidation 1: involved only one soaking step in an aqueous solution of DAP 1 M. Consolidation 2: it was based on a first soaking in 0.5 g/L $\text{Ca}(\text{OH})_2$ NPs dispersion in 2-propanol for 2 hours, followed by a completely drying under hood for one week, then a second soaking was done in an aqueous solution of DAP 1 M for 2 hours, the following two weeks the sample was maintained in a confined environment at RH - 75%. Acronym NT indicates the untreated bone.

Efficacy of the consolidation treatments



Gas porosimetry measurements





Paleogenetic analysis on treated and untreated fragments

Results of the mitochondrial genome sequencing performed on the bones from the Neolithic site of Kierzkowo.

Sample Name	Endogenous DNA (%)	Mean Coverage (X)	Fold Coverage ($\geq 5X$) (%)	DMG 1st Base 3'	DMG 1st Base 5'	Average fragment length (bp)	ContamMix	Haplogroup
K3.4 consolidation 1	18.56	126.52	99.95	0.32	0.33	58.09	0.98	U5b2b1a1
K3.4 consolidation 2	12.52	39.05	99.69	0.30	0.31	61.54	0.98	U5b2b1a1
K3.4 untreat	6.18	14.79	94.27	0.31	0.31	54.69	0.92	U5b2b1a1
K5.3 consolidation 1	0.08	14.94	97.51	0.32	0.38	51.48	0.98	U5b1d1a
K5.3 consolidation 12	5.54	14.84	97.95	0.27	0.29	63.11	0.91	U5b1d1a
K5.3 untreat	5.35	31.76	99.78	0.33	0.30	56.74	0.99	U5b1d1a
K6.2 consolidation 1	3.44	11.72	95.89	0.18	0.17	56.73	0.82	J1c
K6.2 consolidation 2	27.53	60.74	99.38	0.22	0.22	60.1	0.93	J1c
K6.2 untreat	32.58	362.91	99.91	0.22	0.22	60.17	0.97	J1c



Cited papers

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Original article

Evaluation of Diammonium hydrogen phosphate and Ca(OH)₂ nanoparticles for consolidation of ancient bones



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