Sedimentary ancient DNA (sedaDNA) reveal shifts in marine protist communities after World War II and agricultural pollution

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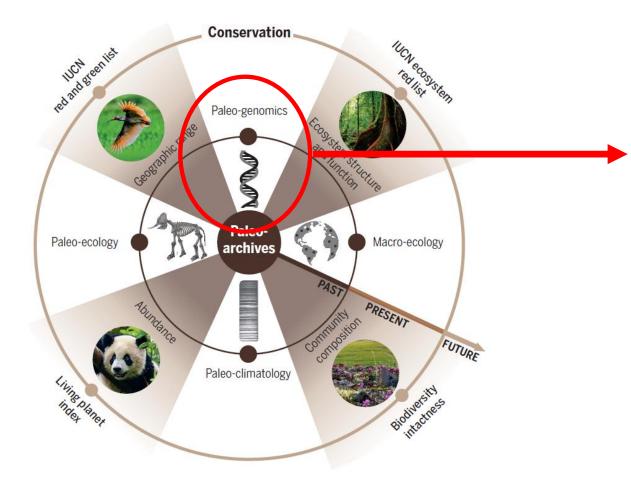






ESTIVAL ROOCH 14-15 Octobre 2021

Paleo-archives: a source of information for understanding ecosystems variations



Using paleo-archives to safeguard biodiversity under climate Fordham et al. 2020 Science Sedimentary ancient DNA



A tool for reconstructing the biodiversity of past biological communities and their shifts in relation to environmental changes.

Plankton Paleogenetics

Ancient Plankton DNA can be preserved in sediments of up to the Pleistocene (~ 125 000 years)

Geobiology

Geobiology (2011), 9, 377-393

DOI: 10.1111/j.1472-4669.2011.00290.x

Preservation potential of ancient plankton DNA in Pleistocene marine sediments

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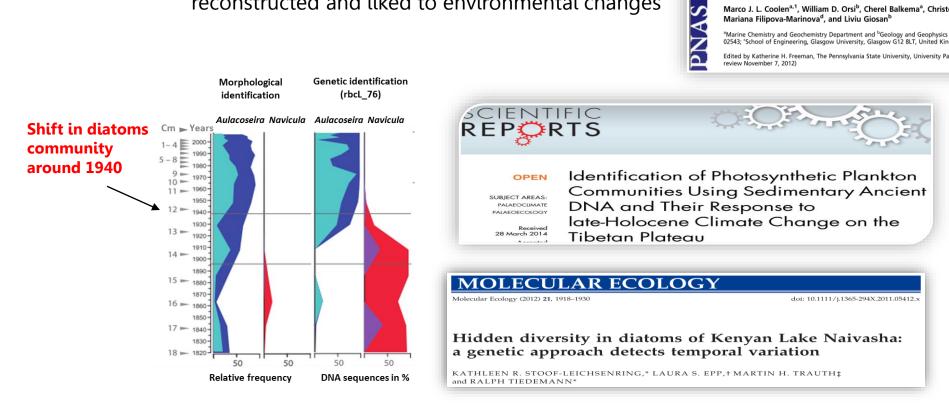
Plankton communities dynamics have been reconstructed and liked to environmental changes

Evolution of the plankton paleome in the Black Sea from the Deglacial to Anthropocene

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Target ecosystems in paleogenetics

Typical deep ocean, permafrost, ice, fjörds, lakes

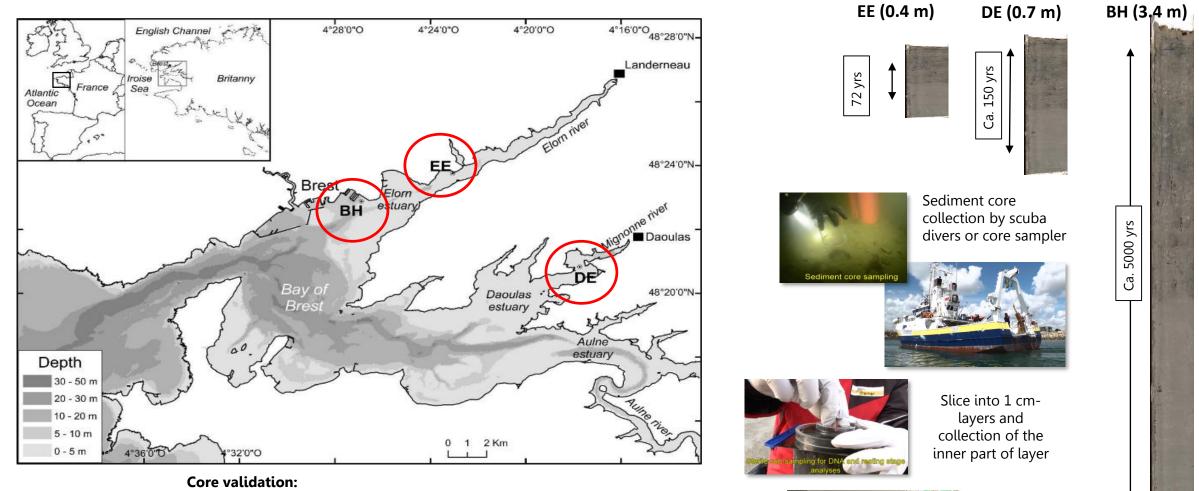
What about coastal ecosystems?

Objectif : Use the potential of paleoecological approach to reconstruct environmental changes of the coastal ecosystem

The Bay of Brest (France)

- 17th -18th centuries: naval town during the industrial développement
- 1940- 1944 : World War II (pollution of army activity, navy traffic, bombing)
- Since 1950 : intense agricultural development (eutrophication)
- Since 1990: control of eutrophication

Sediment core sampling strategy



DNA extraction in dedicated clean lab

- Sedimentation and bioturbation rates (short half-life radionucleotides)
- Sediment dating (²¹⁰Pb, ¹³⁷Cs, ¹⁴C analyses)
- Sediment permeability [H₂0]
- granulometry

Protist sedaDNA nature and its potential use as proxy for ecosystem variation

→ *sed*aDNA degradation (= fragmentation)

Muddy sediments allow a better preservation of the *sed*aDNA that can be analyzed with a good taxonomic resolution (ca. 400 bp marker gene, V4 18 rDNA)

(Capo et al. Mol. Ecol., 2016)

→ sedimentary ancient DNA (*sed*aDNA) characterization :

sedaDNA mostly correspond to intracellular DNA, protected in cell restring stages

(Siano et al. Current Biology, 2021)

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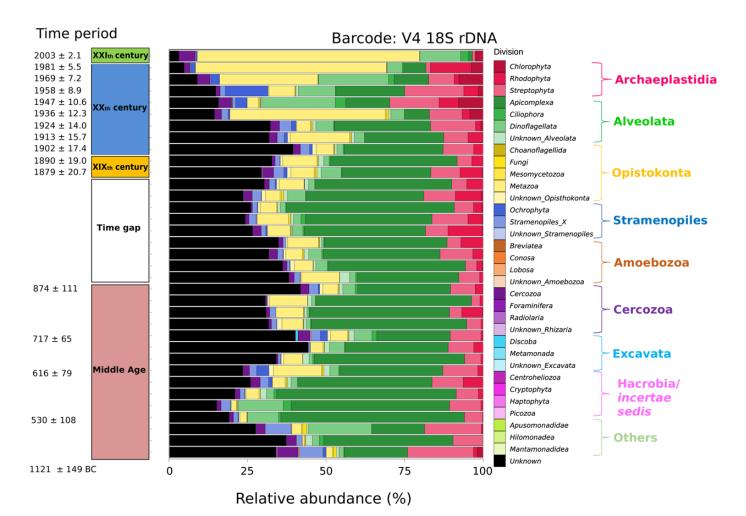
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 sedaDNA as a proxy of protist community shifts in relation to anthropogenic pressures Can sedaDNA contribute to the evaluation of coastal ecosystems resilience? Metacommunity vs. pollutant (heavy metals, PCBs) analyses (Siano et al. Current Biology, 2021)

sedaDNA : protist diversity

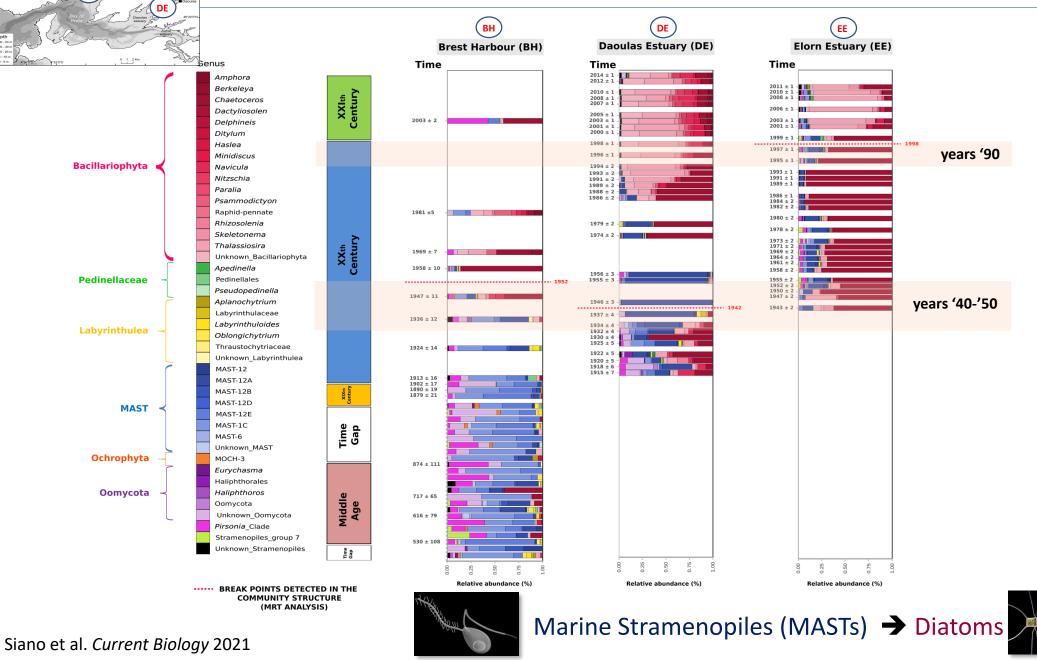


Gregarines (metazoans parasites) very abundant

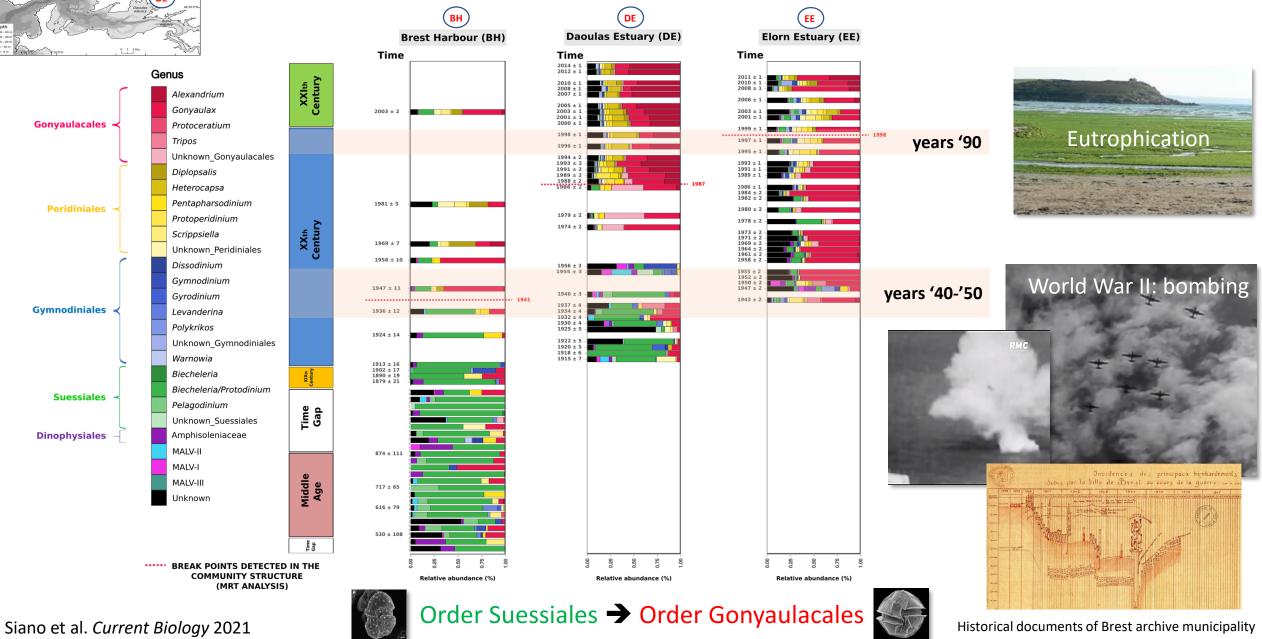
➔ Common pattern with terrestrial protist sediment communities

Siano et al. Current Biology 2021

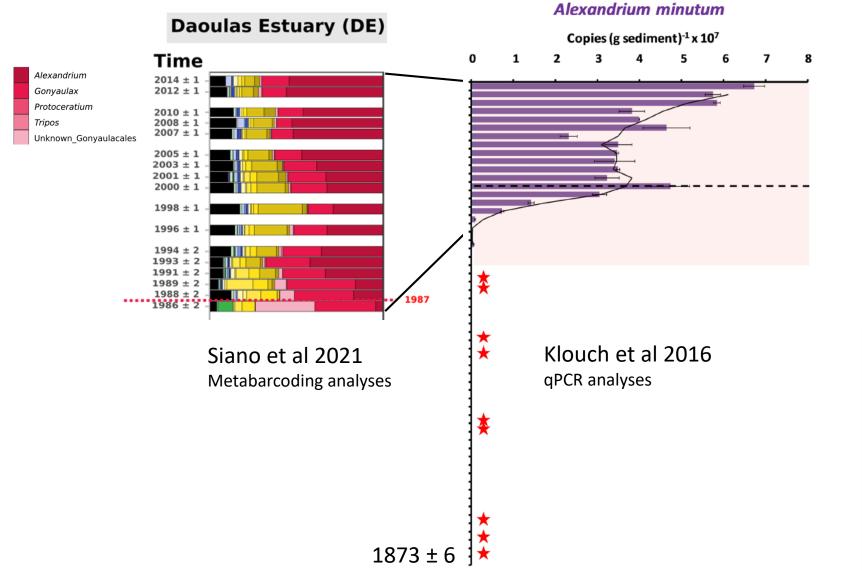
Protist community changes: Stramenopiles



Protist community changes: Dinoflagellates



Dinoflagellate genera shifts





Recrudescence of blooms of the toxic dinoflagellate *Alexandrium minutum*

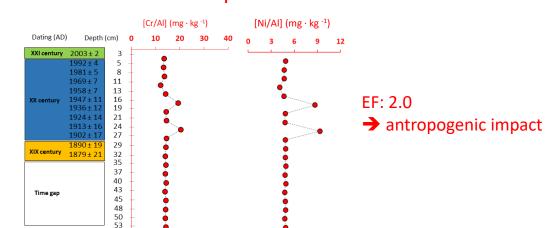


Heavy metal contaminations

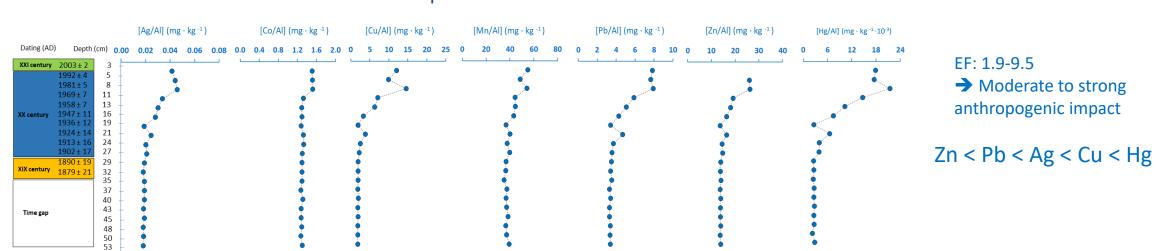
Three heavy metal profile groups and Enrichment Factor Calculation (EF)

[Cd/Al] (mg · kg -1) [V/AI] (mg · kg -1) [Li/Al] (mg · kg -1) Dating (AD) Depth (cm) XXI century 2003 ± 2 1992 ± 4 5 1981 ± 5 8 11 13 1958 ± 7 16 19 21 1913 + 124 27 29 XIX century 1879±2 32 35 37 40 43 Time gap 45 48 50 53

Groupe 1

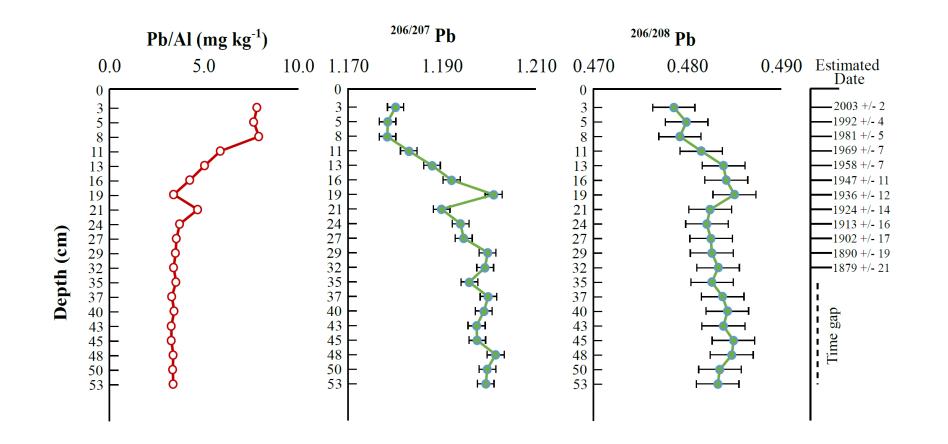


Groupe 2

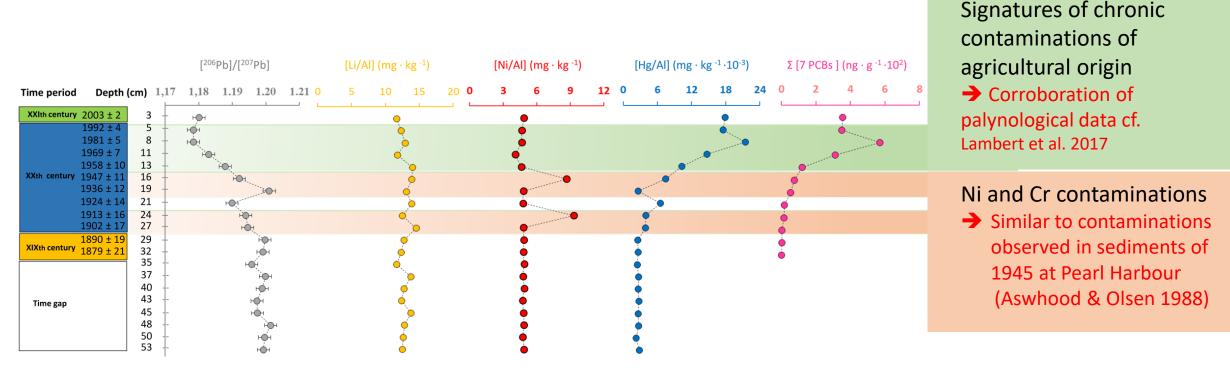


Groupe 3

Heavy metal contaminations

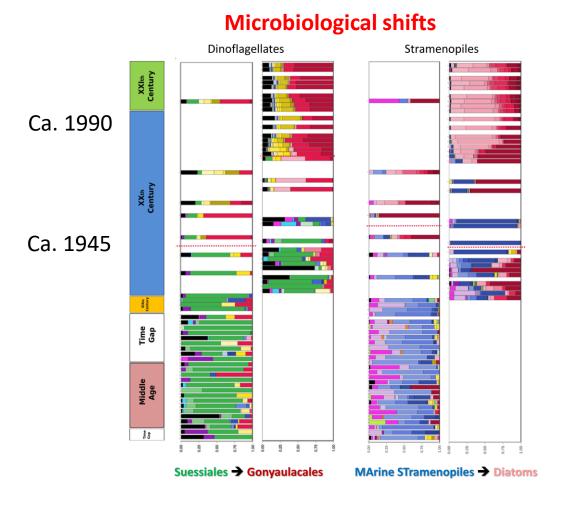


Reconstruction of historical pollutions



Cumulation of occasional (extreme event) and chronic chemical contamination

Conclusions



Contamination



Human impacts







Coastal plankton communities affected by an anthropogenic perturbations were not able to recover their initial (pre industrial) state,

questioning about the resilience and stability capacities of marine coastal areas impacted by humans

Thanks for your attention

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Underwater sediment core sampling image