

# **Soluble Organic Matrix** of axial skeleton and sclerites of Corallium rubrum: partial and comparative analysis of biomineralization



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### Context:

Organic matrix (OM) is the organic fraction located inside biomineral structures of various species.

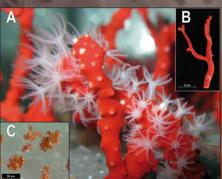
OM is known to have an important role in biomineralization processes. The development of extraction techniques of OM proteins and their characterization thus represents a key step towards an improved understanding of biomineralization.

Comparative studies between different species and models are essential to a greater understanding of biomineralization.

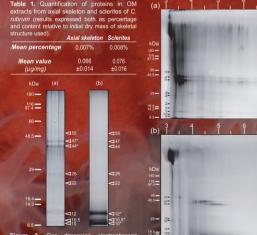
#### Introduction:

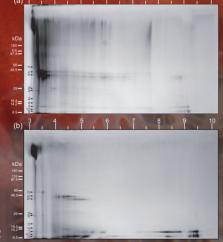
The precious Corallium rubrum (Linnaeus, 1758) from the Mediterranean Sea produces 2 different biominerals: the skeleton and the sclerites (Fig. 1, B, C).

By comparative studies of OM proteins of its 2 skeletal structures, C. rubrum provides an unique opportunity to determine the different and common points in the machinery of the biomineralization process.



## Results:

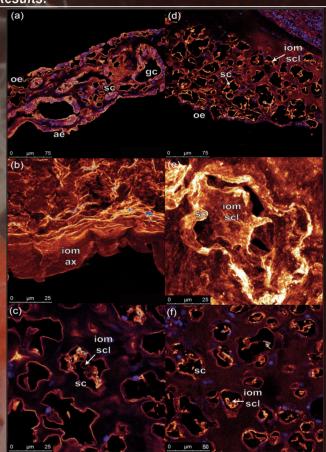




#### Gel electrophoresis analysis:

- 1D gel shows protein bands of similar molecular weight (MW) both for axial skeleton and sclerites. However different intensities are observed (Fig. 2).
- 2. In 2D gels the protein bands are revealed as horizontal lines of protein spots with different Isoelectric points (Fig. 3).
- 2D gels allow better discrimination of OM proteins.

# Results:



#### Organic matrix localization:

- Antibodies against axial skeleton and sclerites both label all tissues with a stronger labeling of the skeletogenic tissues (i.e aboral epithelium and scleroblasts; Fig. 4).
- Decalcified axial skeleton and sclerites are labeled by antibodies and show ghosts of the mineral structures.

# Conclusions and perspectives:

- 1. Axial skeleton and sclerites show similar quantity of OM proteins (Tab. 1) and seem to be composed of a set of OM proteins sharing apparent MW but differing in their biochemical properties (potential post-translational modifications).
- Describing OM proteins opens possibilities for further investigations on sequencing and will be extended to other populations of C. rubrum and other Corallium species.
- Potential use of OM proteins characterization as a tool for:
  - i) Corallium species identification?
  - ii) taxonomic classification?

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