

Understanding ecological functionning of ecological habitats, and building New Indicators based on Genetic Tools to assess their GES (Good Environmental Status).

Aurélien De Jode, Sophie Dubois, Zinovia Erga, Dorian Guillemain, Romain David, Frédéric Zuberer, Jean-Pierre Feral, Anne Chenuil

▶ To cite this version:

Aurélien De Jode, Sophie Dubois, Zinovia Erga, Dorian Guillemain, Romain David, et al.. Understanding ecological functionning of ecological habitats, and building New Indicators based on Genetic Tools to assess their GES (Good Environmental Status).. OT-MED, Feb 2015, Marseille, France. 2015. hal-03663687

HAL Id: hal-03663687 https://hal.science/hal-03663687

Submitted on 10 May 2022

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



Distributed under a Creative Commons Attribution 4.0 International License



Understanding ecological functioning of coralligenous habitats, and building New Indicators based on genetic tools to assess their GES (good environmental status)





Aurélien DE JODE¹, Sophie DUBOIS¹, Zinovia ERGA¹, Romain DAVID¹, Dorian GUILLEMAIN¹, Frédéric ZUBERER², Jean-Pierre FERAL¹, Anne CHENUIL¹, with contributors^{*}

(1) Institut Méditerranéen de Biodiversité et d'Ecologie (IMBE), Station Marine d'Endoume, (2) Service plongée Institut Pytheas aurelien.dejode@imbe.fr



The coralligenous is a typical Mediterranean marine habitat, Paradoxically, unlike Posidonia meadows, it is poorly studied. Like the famous tropical coral reefs, it is a complex biogenic habitat, based on encrusting red algae (Ballesteros 2006).

Abstract	Introduction
The aim of this project is to improve our understanding of its functioning and resilience capacity, and establish biodiversity and connectivity patterns among localities and ecological profiles, providing information for rationalizing the design of Marine Protected Area networks and monitoring methods. To reach this goal, a multidisciplinary approach will be used, combining population genetics, community ecology, microbial ecology and physical oceanography. This requests four tasks, partly using the same data, but different analysis methods. 1- Establishment of the taxonomic composition of the engineer corallinale algae. 2- Establishment of the species composition using metabarcoding (together with	 More precisely, it is defined by the bioconcretions of corallines red algae that grows at dim light conditions. Corallines are the main builders, or enginners species of this ecosystems but their diversity and their ability to colonize new sites is poorly understood. The genus of calcareous red algae Lithophyllum is one of the most important builders in the area of Marseilles, but some marine invertebrates, such as <i>Myriapora truncata</i>, directly contribute to the framework of the habitats because they build their own calcareous skeleton (Hong 1980). However coralligenous habitats are built by various organisms and provide very different ecosystems services from recreational diving to shelter or nurseries for baby fishes and other species.
traditional taxanamy and nhata guadrata validation far aama aamalaa)	llatara gana a una a gualliga a gana la abitata

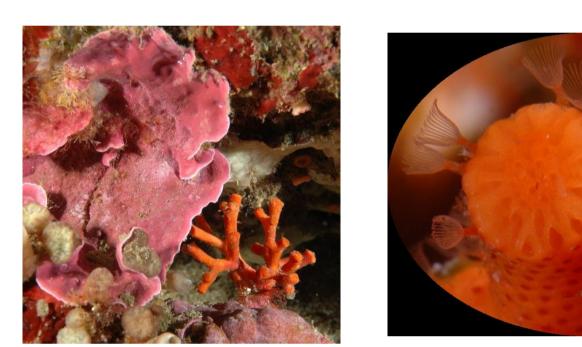
traditional taxonomy and photo quadrate validation for some samples) 3- Population structure and phylogeography of two selected taxa: a red alga *Lithophyllum* spp. and a bryozoan Myriapora truncata, on the CIGESMED sampling network. 4- Synthesis and Comparisons: Propose new GES (good environmental status) indicators for the coralligenous and biodiversity management rules.

Heterogeneous coralligenous habitats

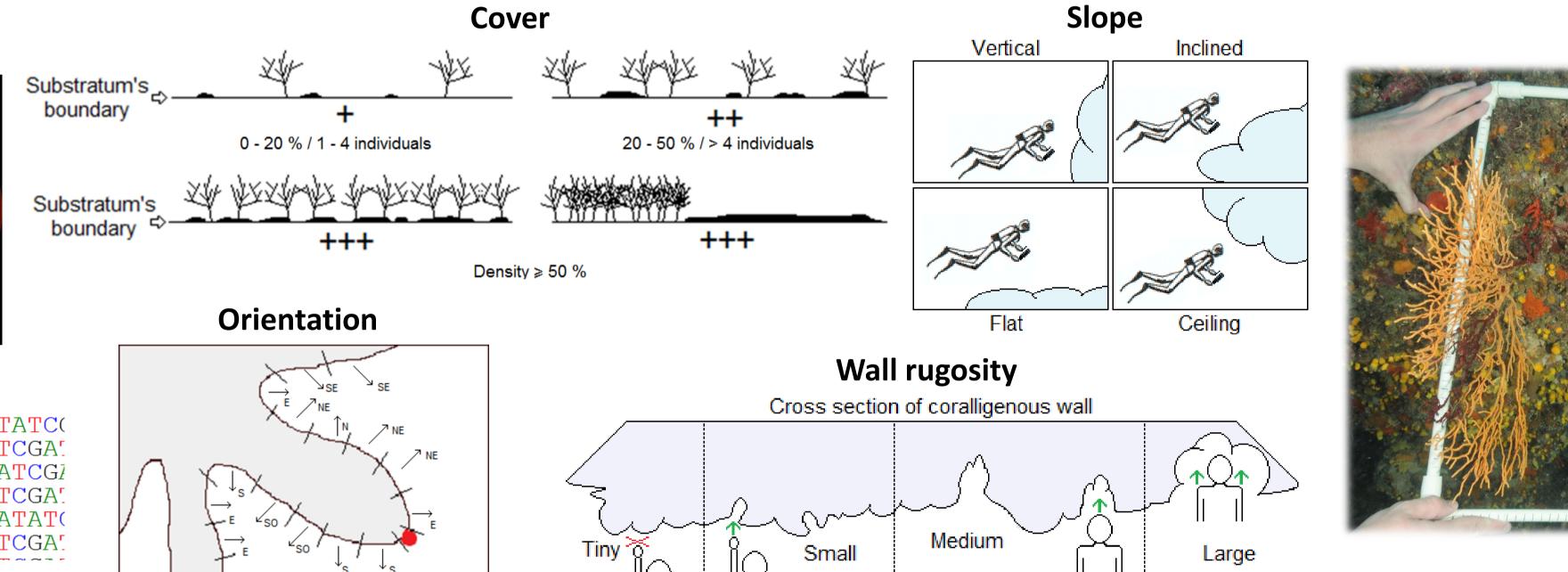


Data

Genetic data obtained after DNA extraction of the samples from the field



I'T'TCGTTGCTGCTCCTCCCGT TCGTTGCTG(Lith CAS 2.5 2 TTGCTTTCGTTGCTGCTCCTCCCGT



Photographic data

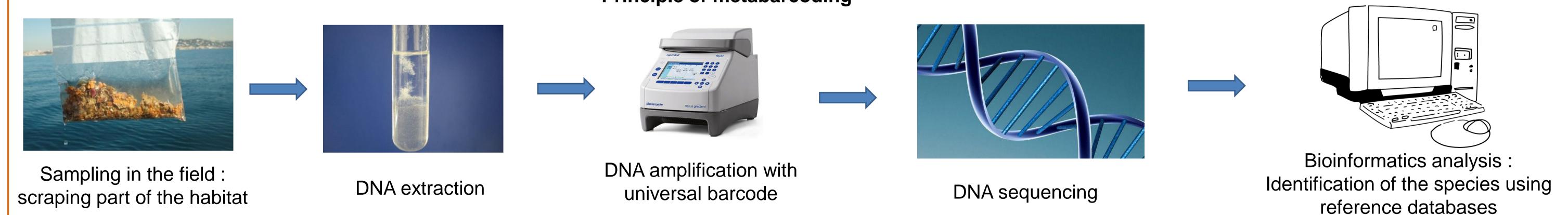


XX

Environmental data directly recorded on the field during the sampling

Method

In this project different methods DNA extraction, PCR, metabarcoding and disciplines will be used : population genetics, phylogeography, community ecology and hydrological modeling. **Principle of metabarcoding**

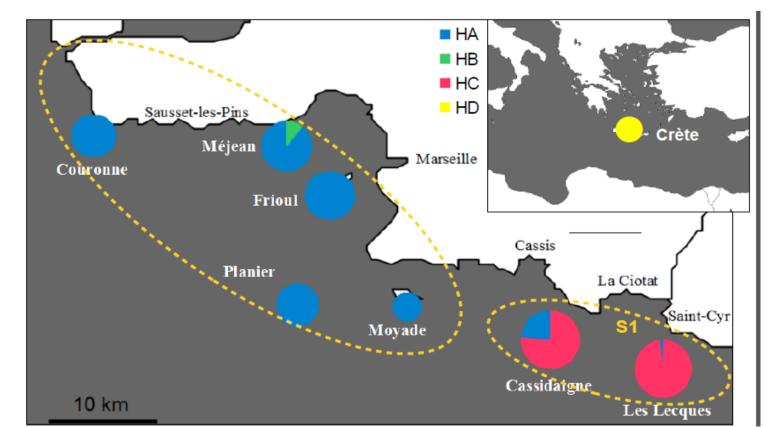


Results

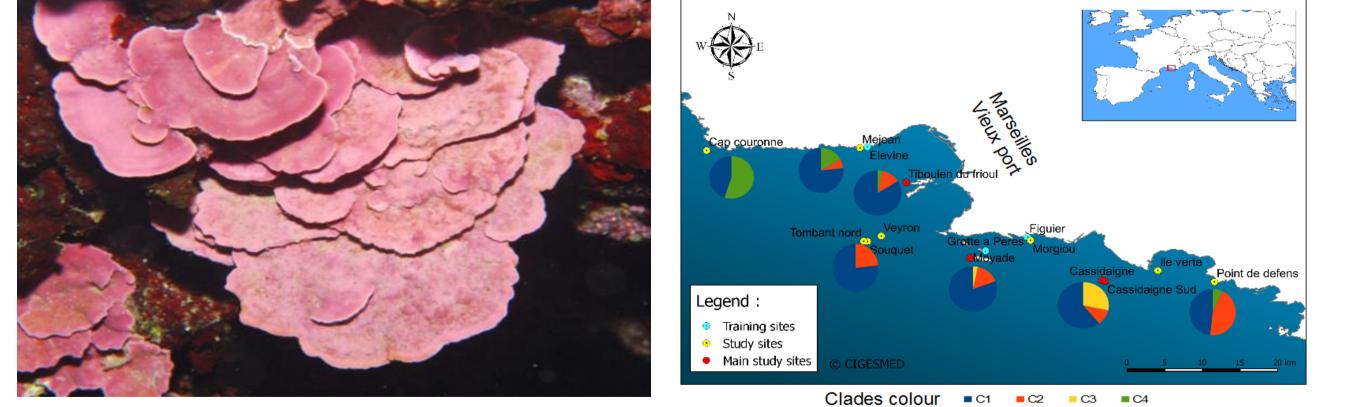
Preliminary studies of the two engineers species revealed genetic differentiation in the Bay of Marseille.

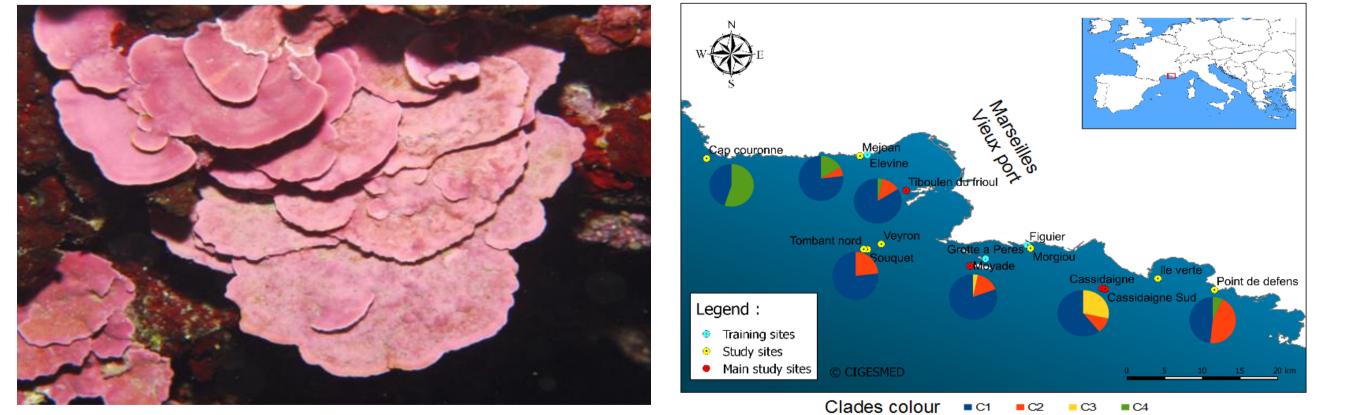
Myriapora truncata











Conclusions

Our first results suggests that the engineers species are actually composed of cryptic species that may have live in different ecological niches. It emphasizes our lack of knowledge about this endemic Mediterranean ecosystem and stresses our requirement for a better understanding of its functioning in order to rationalize its monitoring and protection.

Acknowledgements

This work is a contribution to the Labex OT-Med (n° ANR-11-LABX-0061) funded by the French Government «Investissements d'Avenir» program of the French National Research Agency (ANR) through the A*MIDEX project (n° ANR-11-IDEX-0001-02).

*Other Contributors to the CIGESMED program : Acik Cinar S., Andral B., Arvanitidis C., Aurelle D., Aysel V., Bakir K., Bellan-Santini D., Bouchoucha M., Cahill A., Celik C., Chatzigeorgiou G., Chatzinikolaou E., Chenesseau S., Çinar M.E., Dağli E., Dailianis T., DimitriadisC., D'Iribarne C., Doğan A., Faulwetter S., Gatti G., Gerovasileiou V., Güçver S.M., Issaris Y., Katağan T., Keklikoglou K., Kirkim F., Koçak F., Koutsoubas D., Marschal C., Önen M., Önen S., Öztürk B., Panayiotidis P., Pergent G., Pergent G., Pergent G., Pergent G., Rocher C., Ruitton S., Sakher S., Salomidi M., Sarropoulou E., Sartoretto S., Selva M., Sini M., Sourbes L., Simboura N., Taşkin E., Thierry de Ville d'Avray L., Tschanz L., Vacelet J., Valavanis V., Vasileiadou A., Verlaque M.