LifeWatchGreece Hellenic Centre for Marine Research





- 400 CPU cores & 20 nodes
 up to 640GB RAM on a single node
- 40Gbps Infiniband interconnection



more than 200 state-of-the-art software suites for research analyses



more than 50 users, 20 countries and 30 projects



NGS data analyses (de novo assembly, annotation, SNP discovery, phylogeny, statistics and more)









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- DNA (genomics, genetics, metagenomics, phylogenomics, metabarcoding, population genomics)

- Area 52 lab metabarcoding (Haris Zafeiropoulos, IMBBC, HCMR) View
- Microbial community diversity in the lagoonal complex of Amvrakikos Gulf (Ionian Sea, Western Greece) (Dr. Christina Pavloudi, IMMBC, HCMR) View
- Eukaryotic Microbial Plankton in the Western Levantine Basin (Eastern Mediterranean Sea) (Dr. Iliana Santi, IO, HCMR) View
- High nutrient addition of different intensities alters plankton composition and the biotic relationships among plankton species (Dr. Iliana Santi, IO, HCMR) View
- SPELMED: Population Genomics analysis of the small pelagic fish stocks in the Northwestern Mediterranean Sea (Dr. Costas Tsigenopoulos) View
- Genome analysis of Pseudomonas strains isolated from Kolumbo, Santorini. Panos Bravakos View
- · Hybrid assembly and genome analysis of an epitheliocystis related agent. Maria Chiara Cascarano Phd student View
- Phylogenomics investigation of sparids (Teleostei: Spariformes) using high-quality proteomes highlights the importance of taxon sampling
 (Dr. Tereza Manousaki) View
- . MOUNT: Seascape genomics of invertebrates in coastal Crete View



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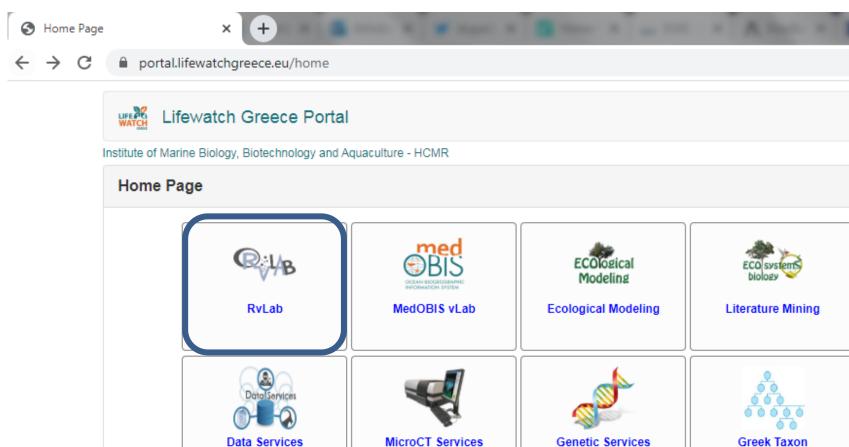
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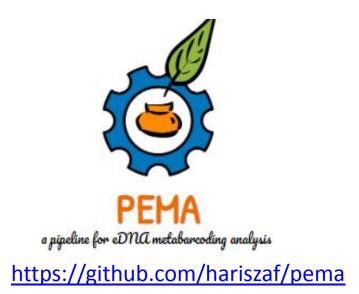


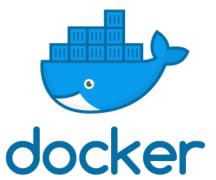






Containerization







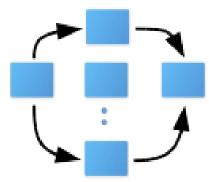
https://singularityhub.org/collections/2295



https://hub.docker.com/r/hariszaf/pema

Containerization







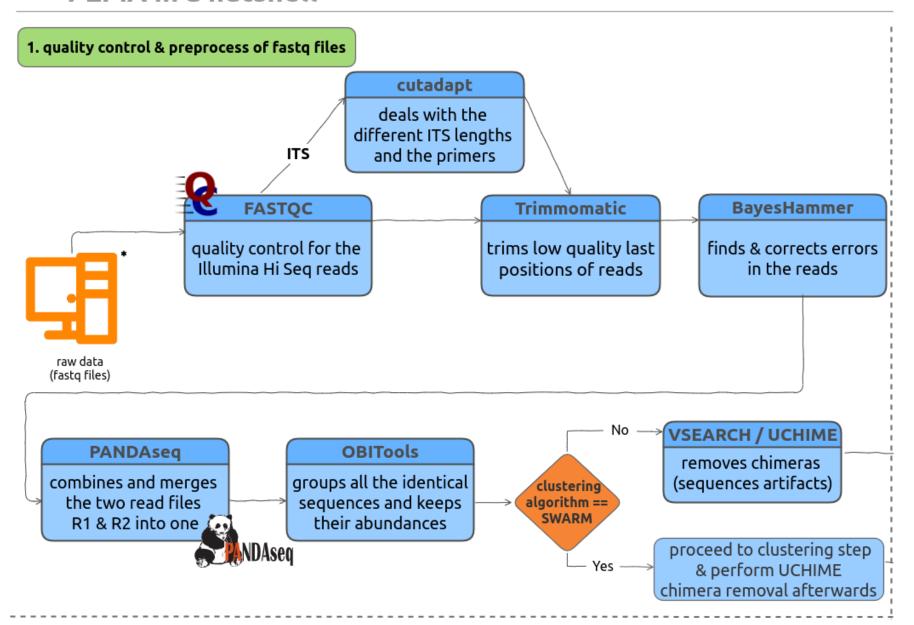
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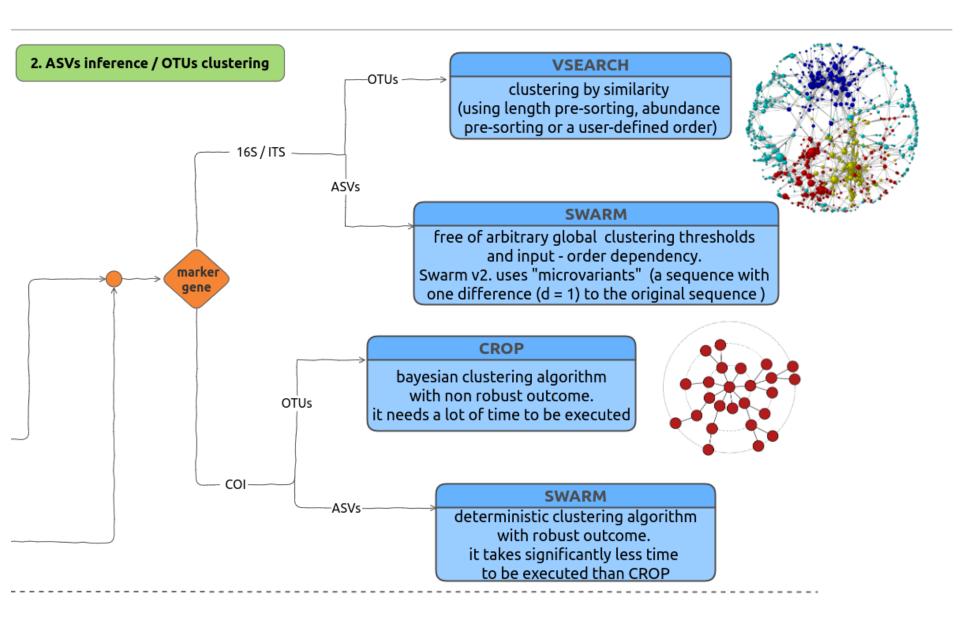




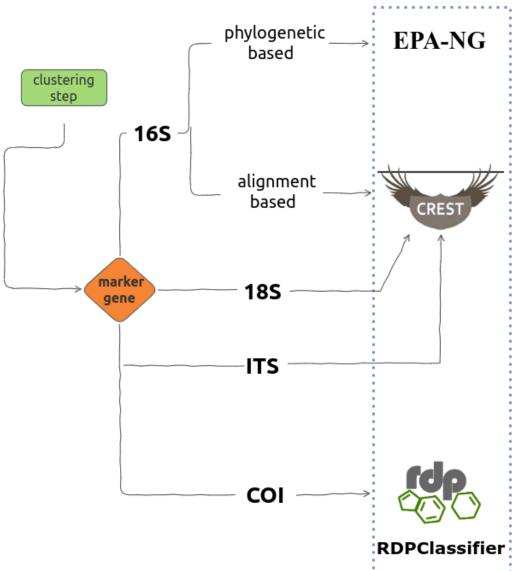
https://hub.docker.com/r/hariszaf/pema

PEMA in a nutshell





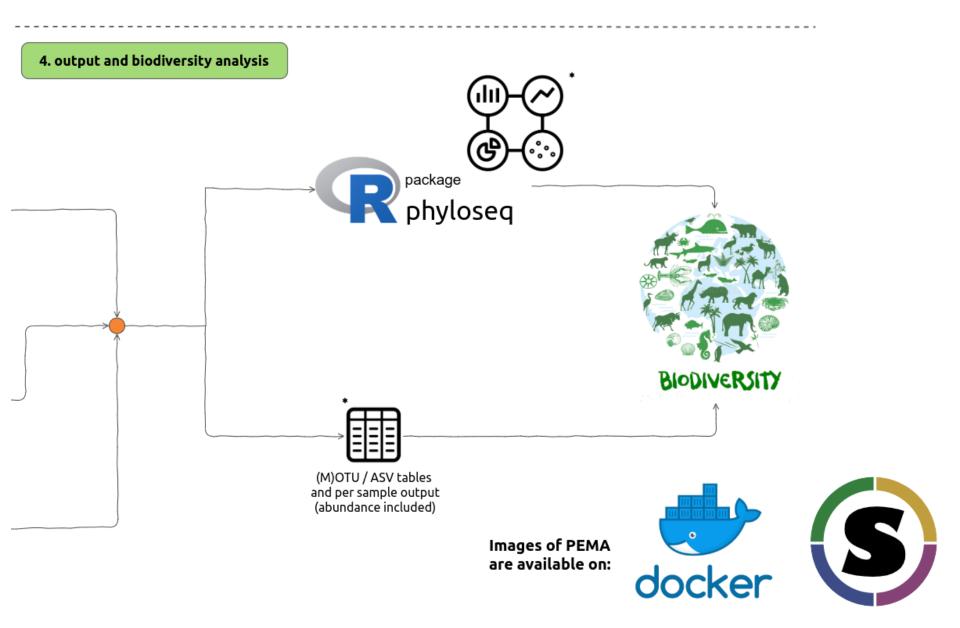
3. methods for taxonomic assignment







+ MIDORI Reference



Artificial Reef Monitoring Structures (ARMS)

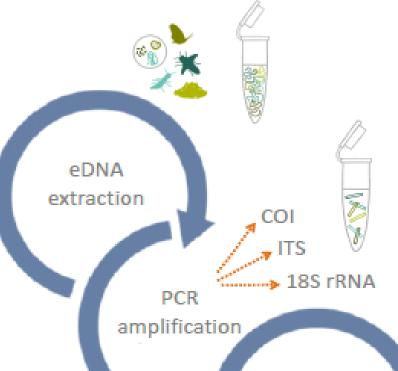


ARMS assembled (before deployment)





ARMS deployed in one of the chosen locations



MiSeq Sequencing

tttgagtatacaact

ttcgagcatacgact
aacgtccaaaggagt
ttggagcatacgact
aaggtccaaagagt
ttcgagcatacgact
atcgtccaatggagt

aaggtccaaacgagt aacgtccaaaggagt

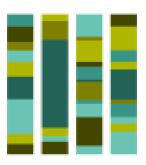
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P.E.M.A.:

a Pipeline for Environmental DNA Metabarcoding Analysis for the 16S and COI marker genes

Sequence analysis



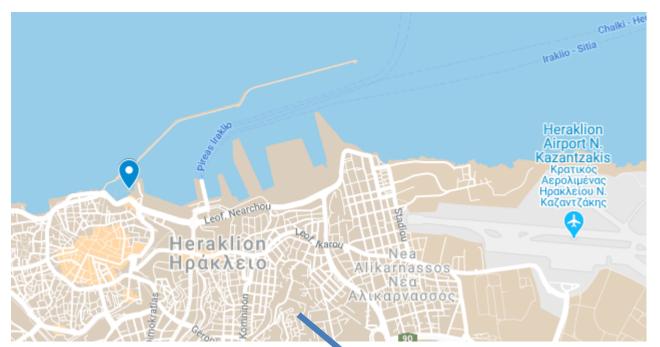
Success story!

First record of the nudibranch *Anteaeolidiella lurana* (Ev. Marcus & Er. Marcus, 1967) in Greek waters

Vasilis GEROVASILEIOU 1, Christina PAVLOUDI 1, Francisco SEDANO 1,2

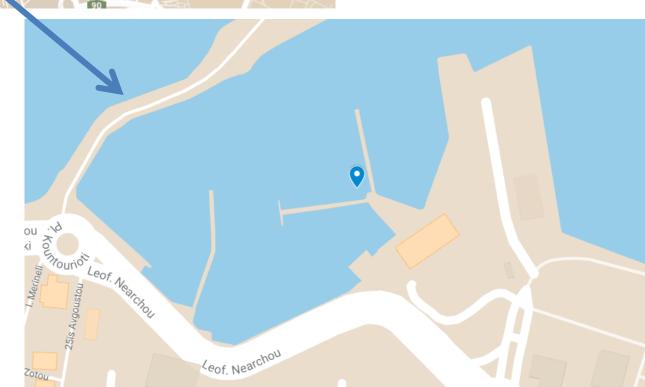
¹ Institute of Marine Biology, Biotechnology and Aquaculture, Hellenic Centre for Marine Research, Heraklion, Greece

² Laboratorio de Biología Marina, Departamento de Zoología, Universidad de Sevilla, Spain



1st deployment: 28/9/2018-28/1/2019

2nd deployment: 12/6/2019-11/10/2019



- 1st deployment: 28/9/2018-28/1/2019
- molecular evidence (eDNA) for the presence of Anteaeolidiella lurana
- \rightarrow How can we be sure?
- Only three of the accepted *Anteaeolidiella* species (*A. indica, A. ireneae,* and *A. orientalis*) do not have sequence data associated with them.

All the rest (8) species, including *A. lurana*, have COI sequences in the **MIDORI reference database**, which was used for the taxonomic assignment.

The degree of certainty for the identification based on the eDNA is high.

2nd deployment: 12/6/2019-11/10/2019

→ photographic documentation



thank you for finding me.

Christina Pavloudi cpavloud@hcmr.gr

Haris Zafeiropoulos haris-zaf@hcmr.gr