An underwater photograph showing a dense bed of mussels on a dark seabed. The mussels are primarily yellowish-brown with some darker ones. Numerous small, translucent shrimp-like crustaceans are scattered throughout the mussel bed and the surrounding water. The water is clear and blue, with some light rays visible. The overall scene depicts a rich marine ecosystem.

Marine Genetic Resources (MGR)
why are they important?
How are they impacted?

Ester Serrão
CCMAR and University of Algarve, Portugal

Marine Genetic Resources (MGR)

Genetic information (DNA) from marine organisms with
current or **potential** value.

direct economic value, ecosystem services, adaptability.

Potential value cannot be determined
future conditions and technologies
unknown

All genetic variation is a resource.

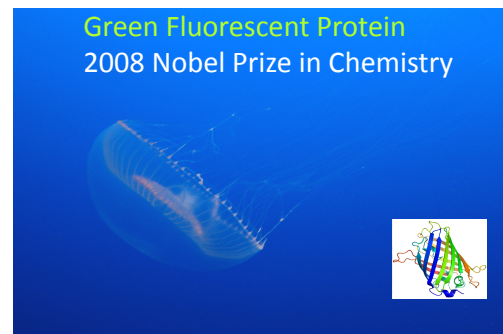
Why conserve MGR?

- Ecosystem services

(marine food webs, biogeochemical cycles, climate)

- Direct uses

(blue biotechnology, fisheries, aquaculture)



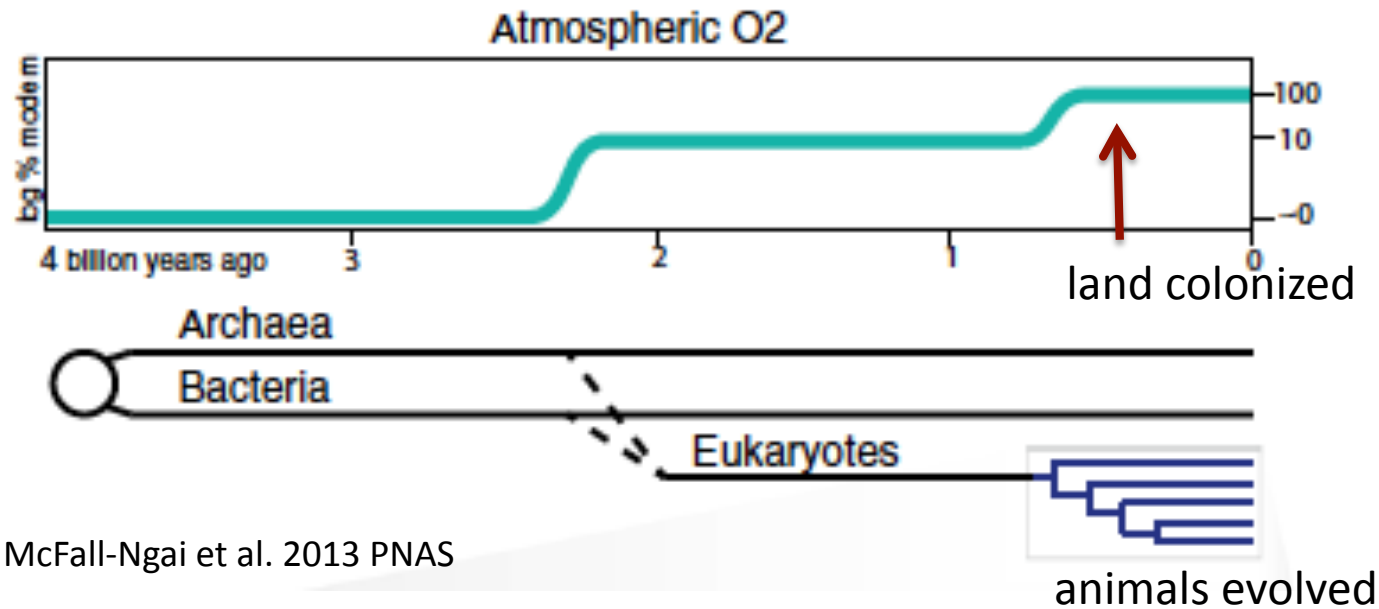
- Adaptation

to continue providing services/uses

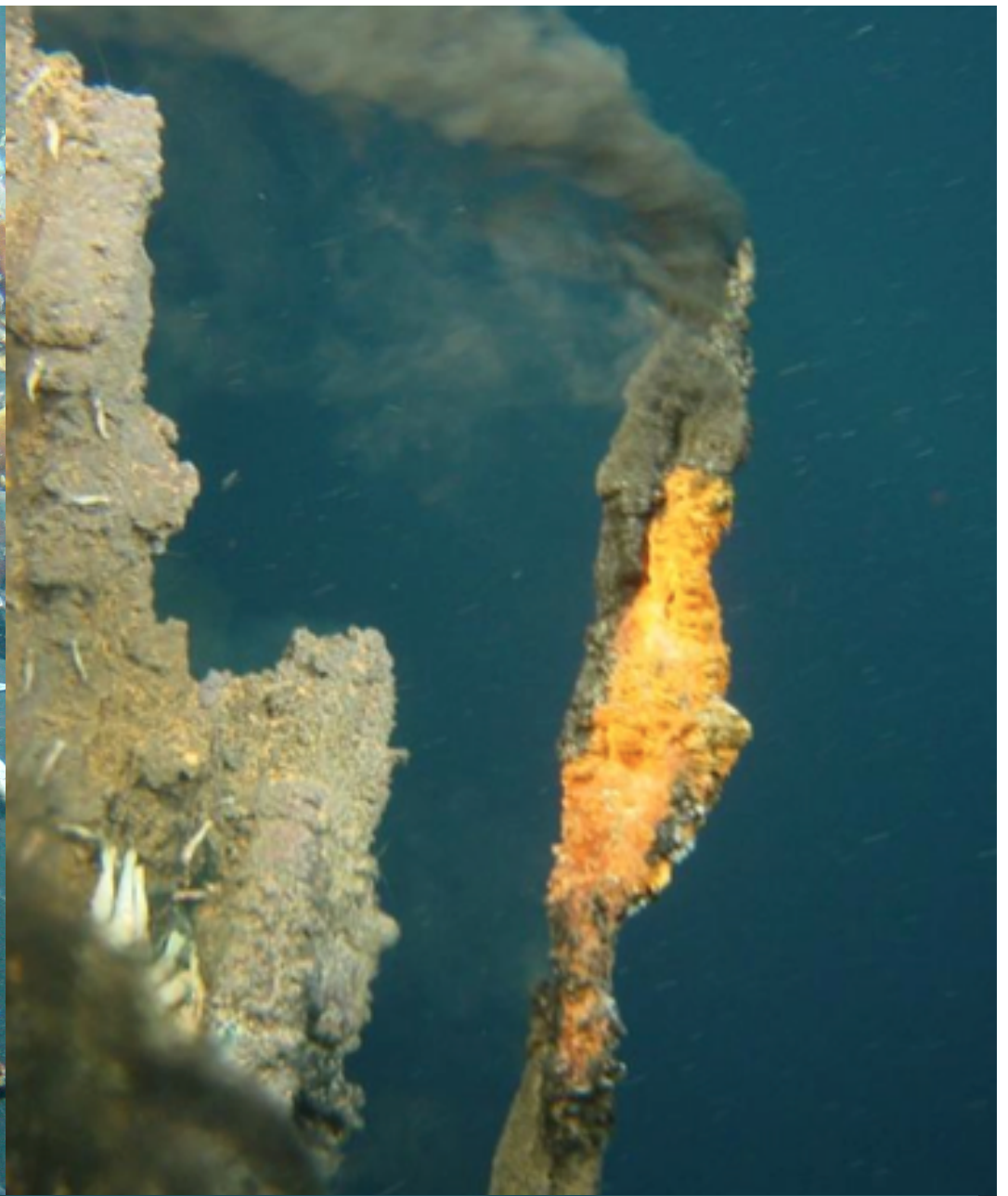
MGR in the ecosystem, billions of years of evolution

Life evolved in the sea mostly as micro-organisms, until ≈ 450 mya when land was colonized

Sea: much more diversity of life forms and processes, more genetic diversity, more biochemical diversity.



Marine microbes have defined the chemistry of the oceans and atmosphere

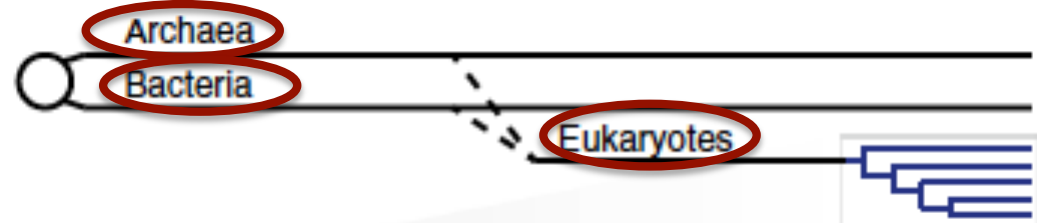
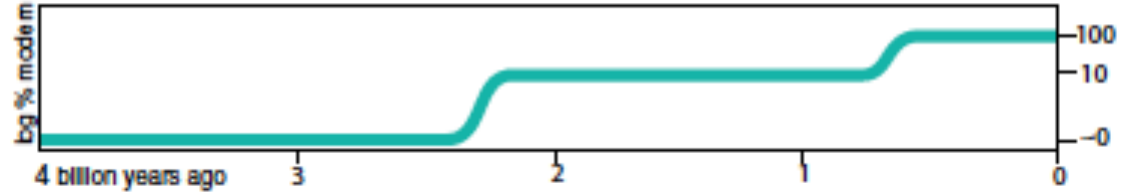


chemoautotrophic microorganisms that can live in symbiosis with invertebrates.

3 Domains of Life

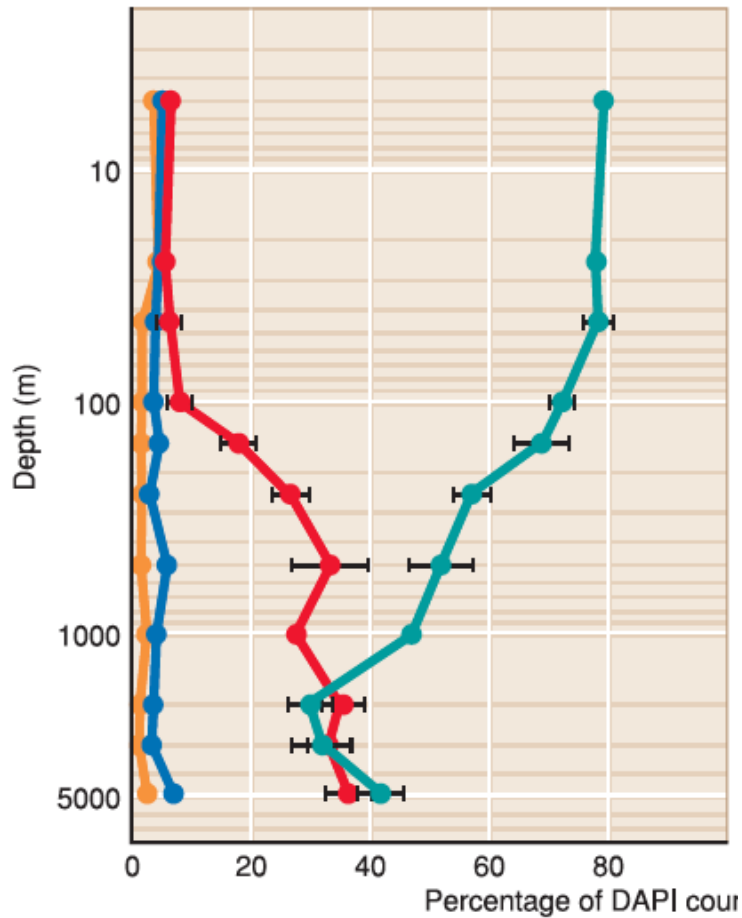
McFall-Ngai et al. 2013 PNAS

Atmospheric O₂



Within the 30 phyla of **animals**, 29 occur in the oceans, 15 are exclusively marine

Microbial domains



- Crenarchaeota
- Bacteria
- Euryarchaeota
- Negative control

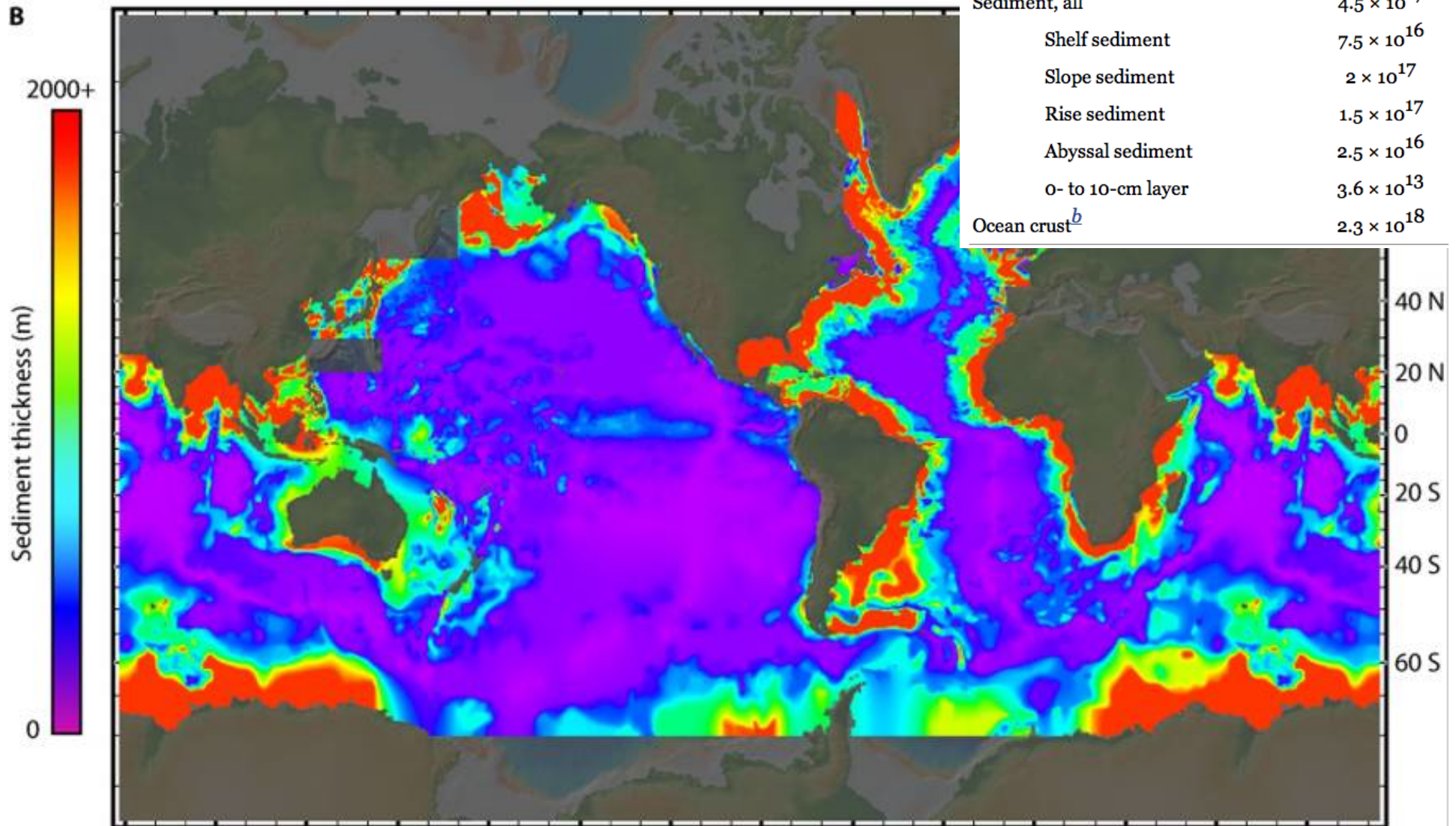
Archaea dominance in the mesopelagic Pacific

Karner, M., E. F. DeLong, and D. M. Karl. 2001. Nature 409:507–510

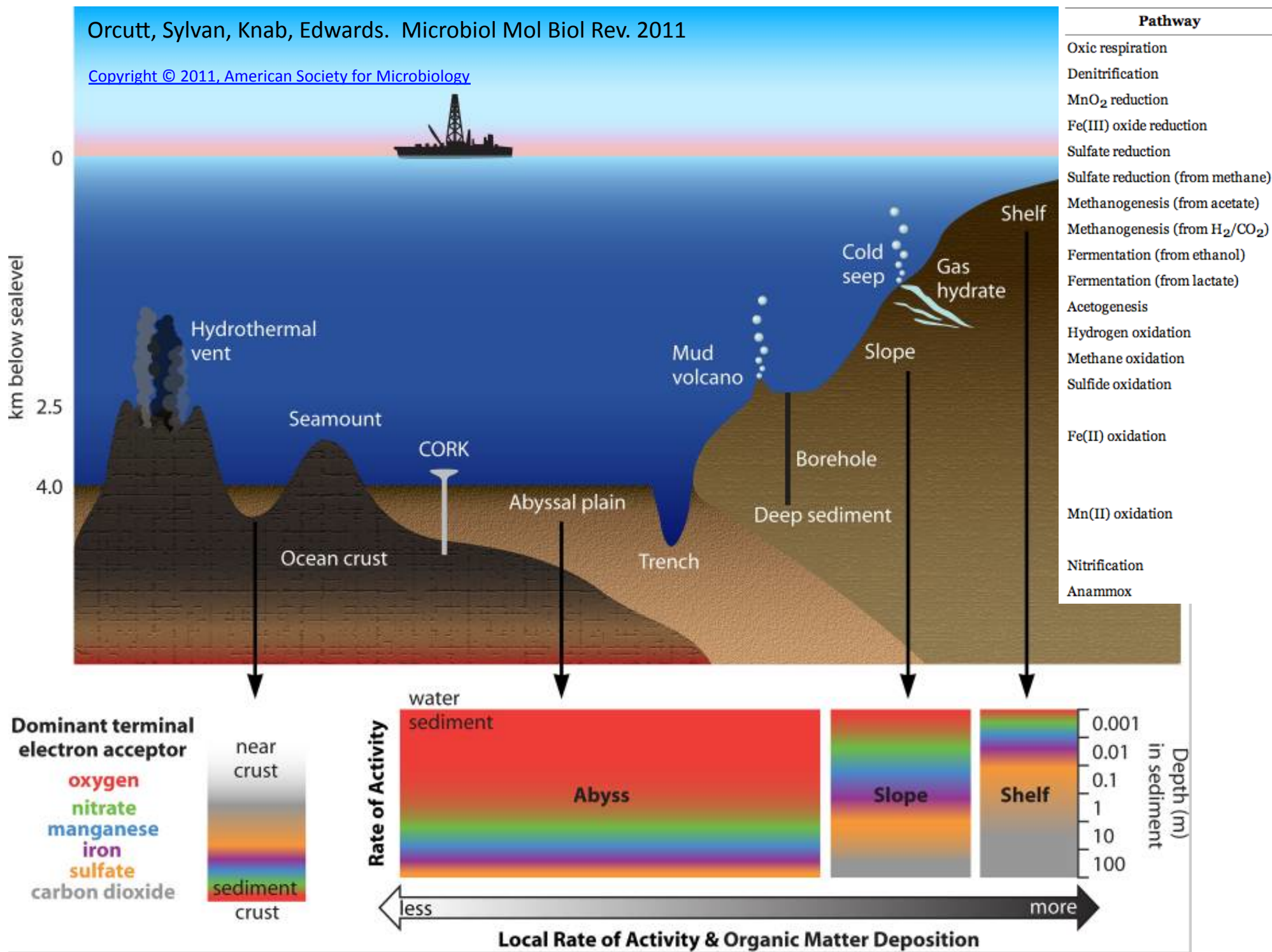
The largest habitats on Earth

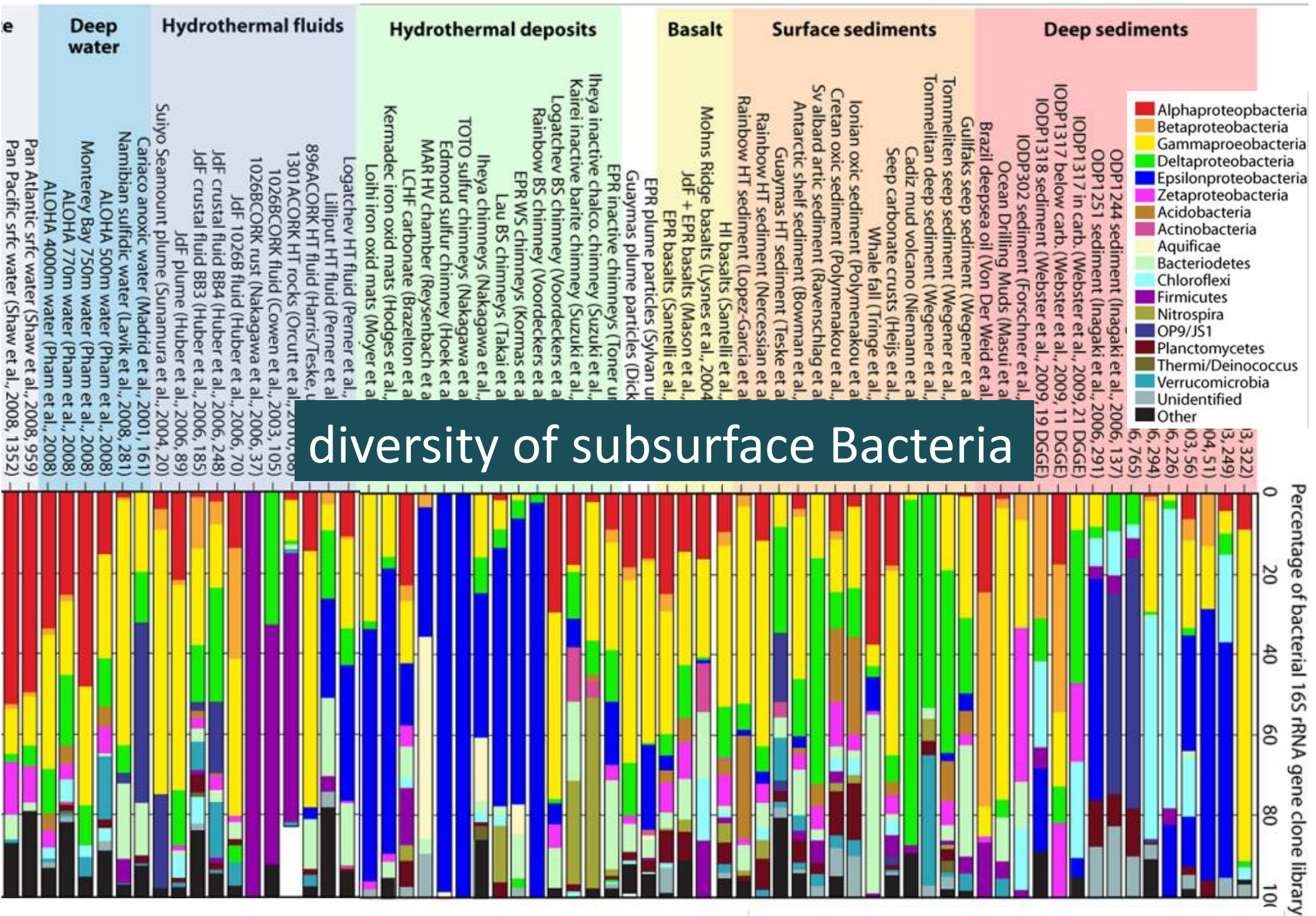
Estimated volumes of various habitats in the dark ocean, and of the photic zone

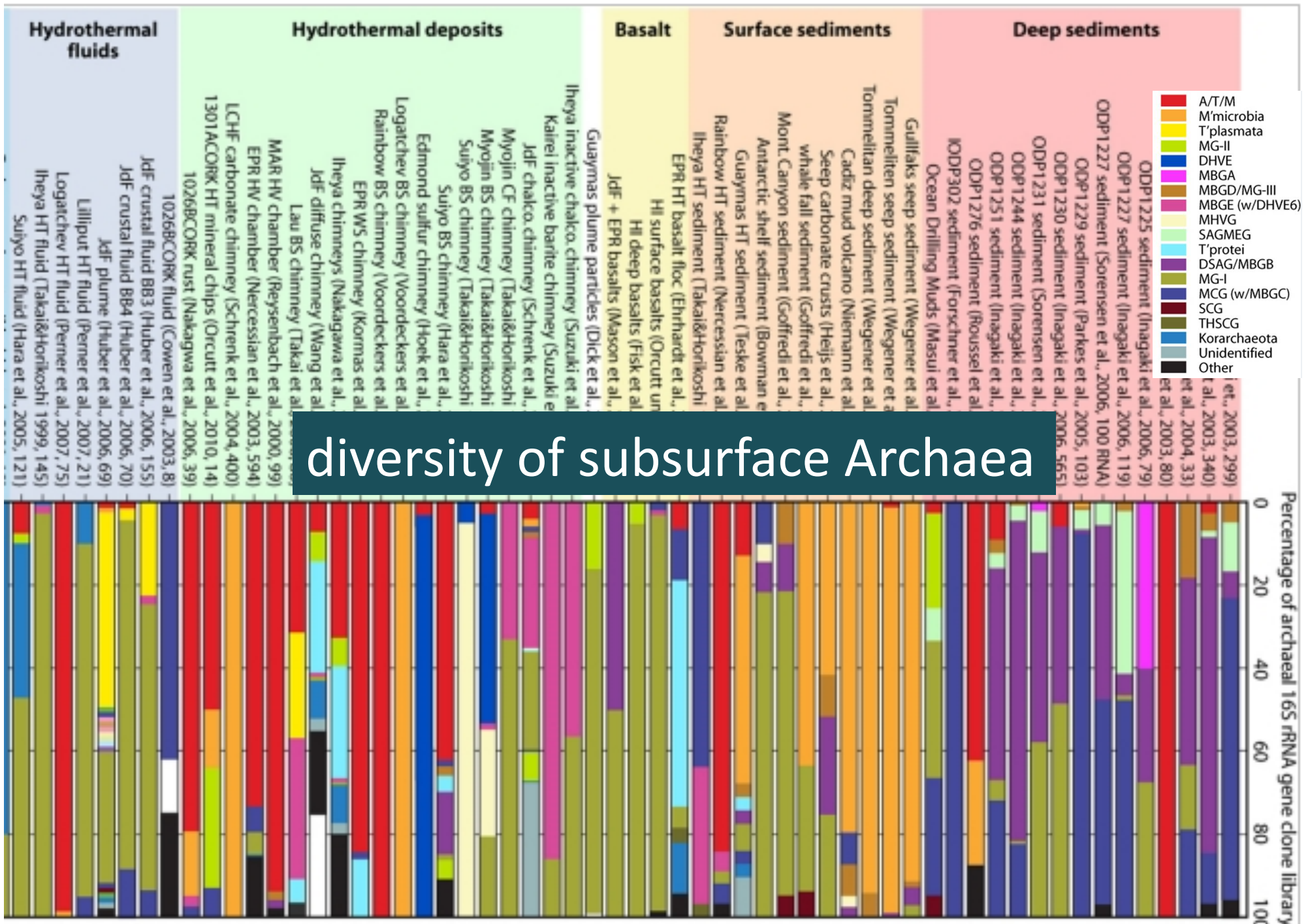
| Habitat | Vol (m ³) |
|---------------------------------------|----------------------------|
| Water column (<200 m below sea level) | 3.0×10^{16} |
| Water column (200+ m below sea level) | 1.3×10^{18} |
| Hydrothermal plumes ^a | 7.2×10^{13} (/yr) |
| Subsurface ocean | $\sim 10^{16}$ |
| Sediment, all | 4.5×10^{17} |
| Shelf sediment | 7.5×10^{16} |
| Slope sediment | 2×10^{17} |
| Rise sediment | 1.5×10^{17} |
| Abyssal sediment | 2.5×10^{16} |
| 0- to 10-cm layer | 3.6×10^{13} |
| Ocean crust ^b | 2.3×10^{18} |



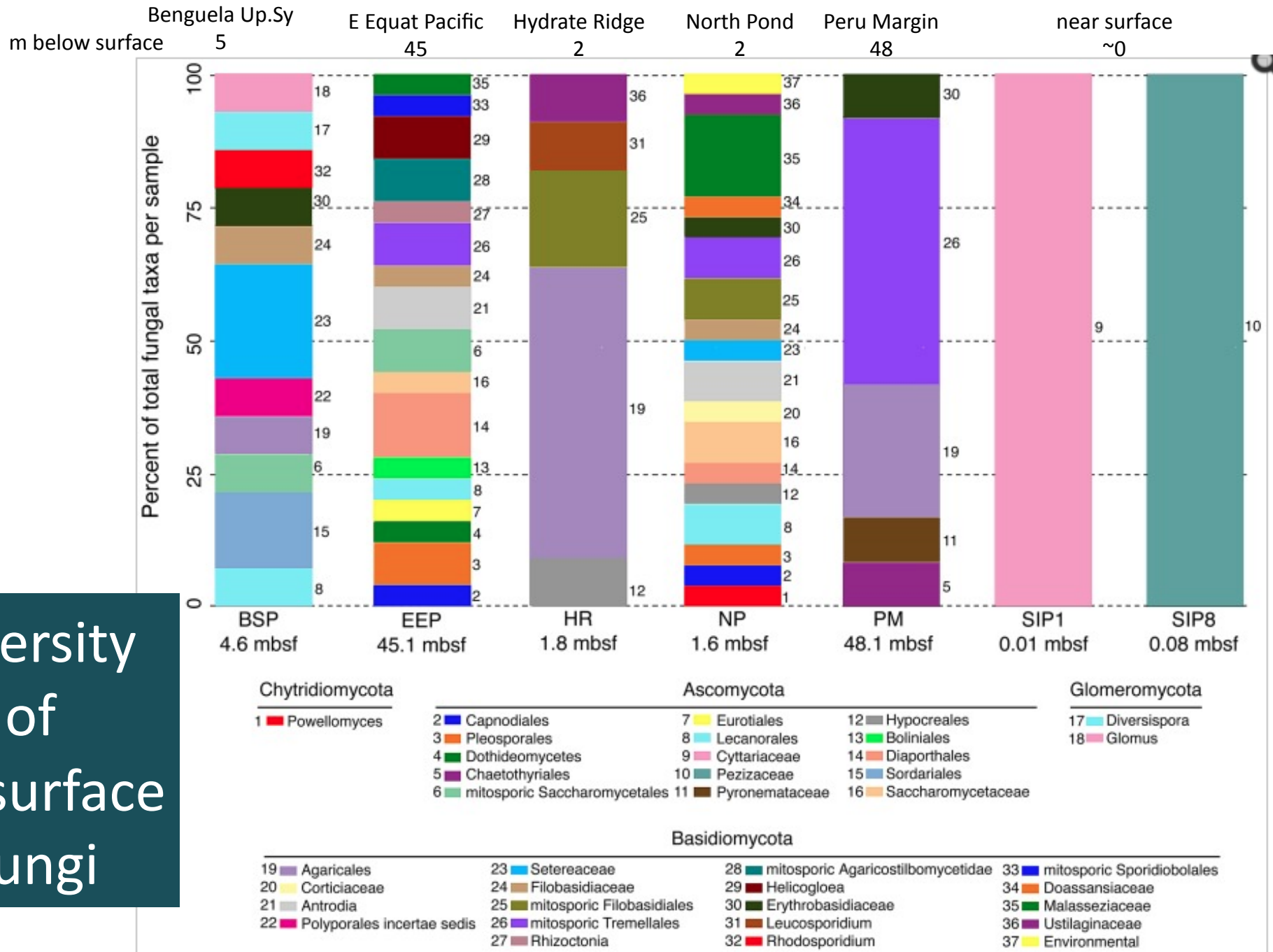
Copyright © 2011, American Society for Microbiology







Deep sequencing of seafloor eukaryotic rRNA reveals active Fungi across marine subsurface provinces.



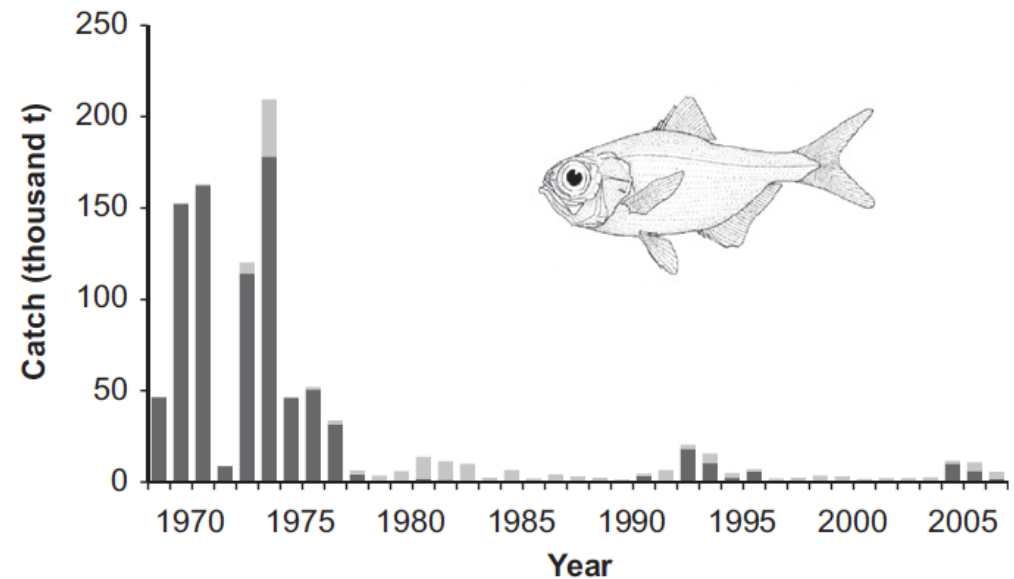
diversity
of
subsurface
Fungi

Impacts on MGR

-reducing **population size** - genetic drift, bottlenecks

-preventing **reproduction** of the majority of the individuals (inbreeding)

demographic collapse → genetic loss



Norse, Brooke, Cheung, Clark, Ekeland, Froese, Gjerde, Haedrich, Heppell, Morato, Morgan, Pauly, Sumaila, Watson. Sustainability of deep-sea fisheries. Marine Policy. 2012.

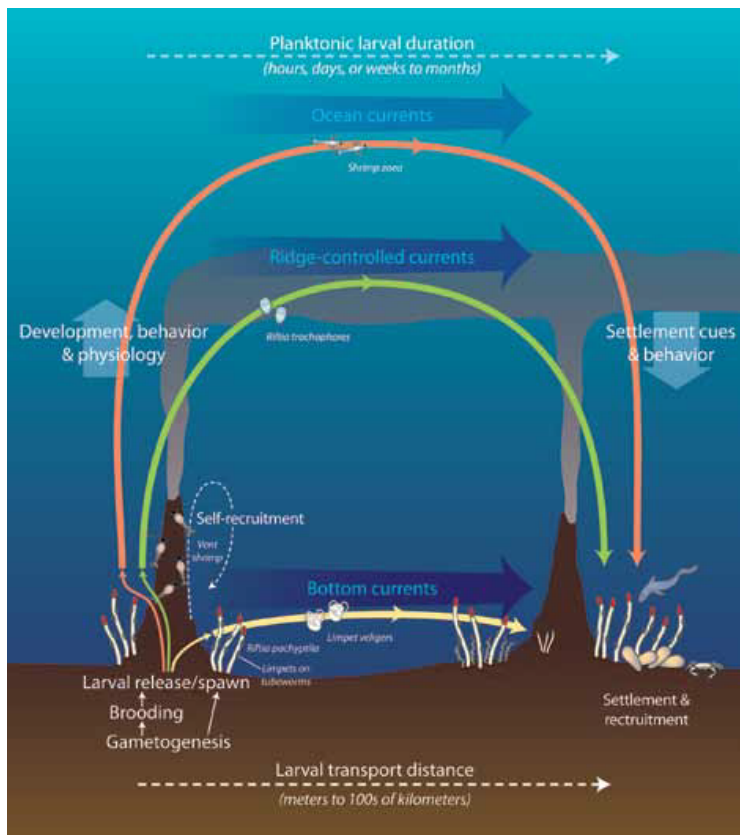
Fig. 4. Catches of slender armourhead (dark gray) and alfonsino (light gray) from Emperor and Hawaiian seamounts [80,133,148]. Splendid alfonsino image: wikipedia.org.

Impacts on MGR

Destruction of habitats:

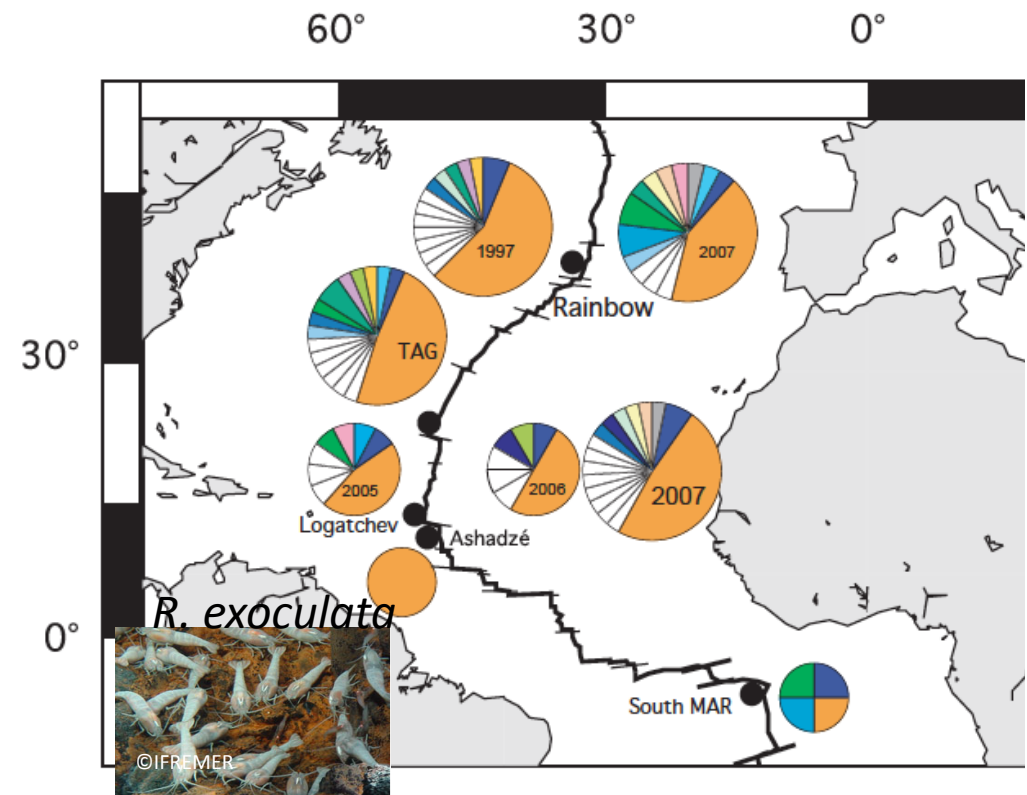
- extinction of genetically distinct or locally adapted populations
- along dispersal routes - gene flow disturbance

Adams, Arellano, Govenar. 2012. Oceanography.



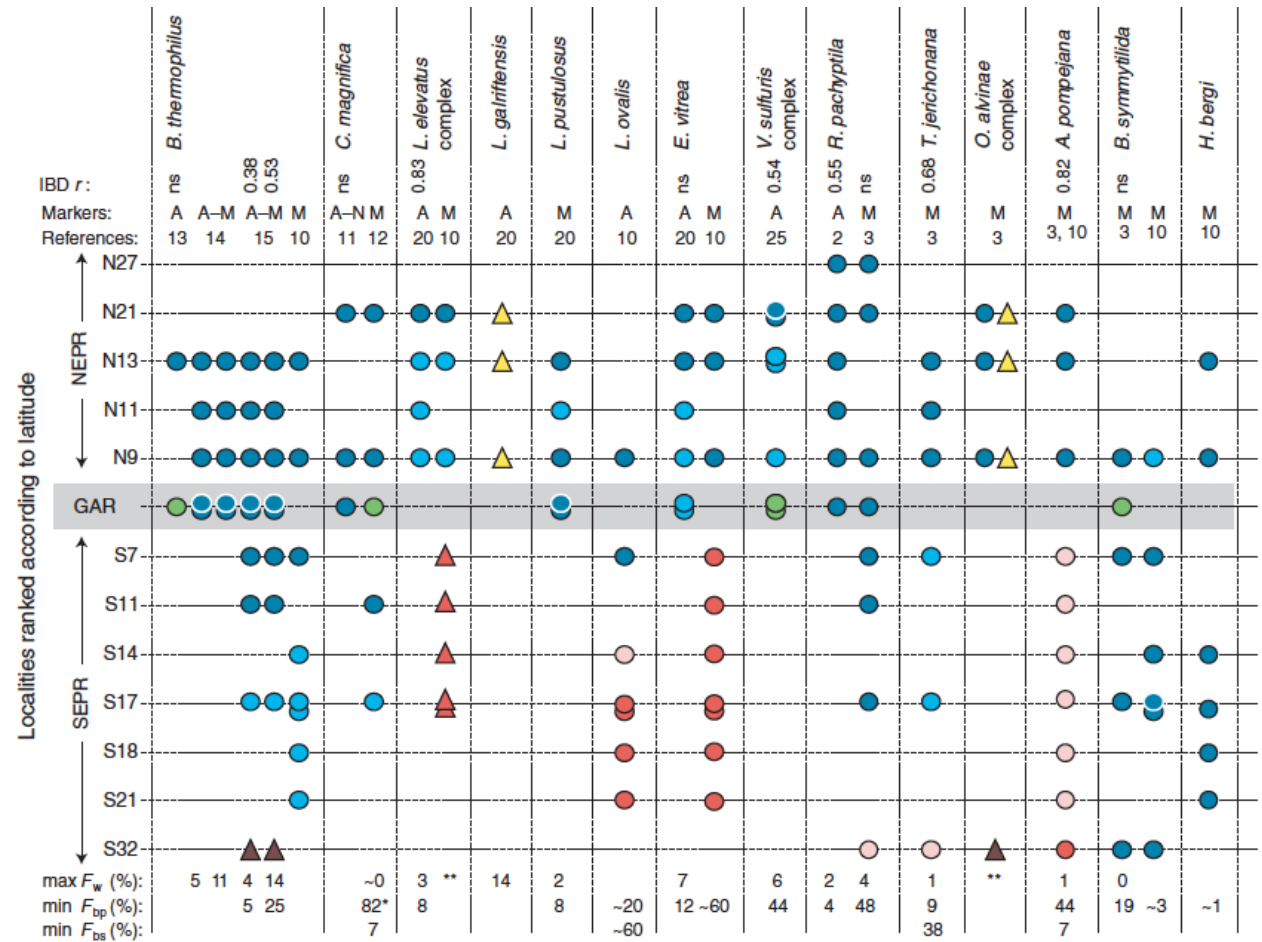
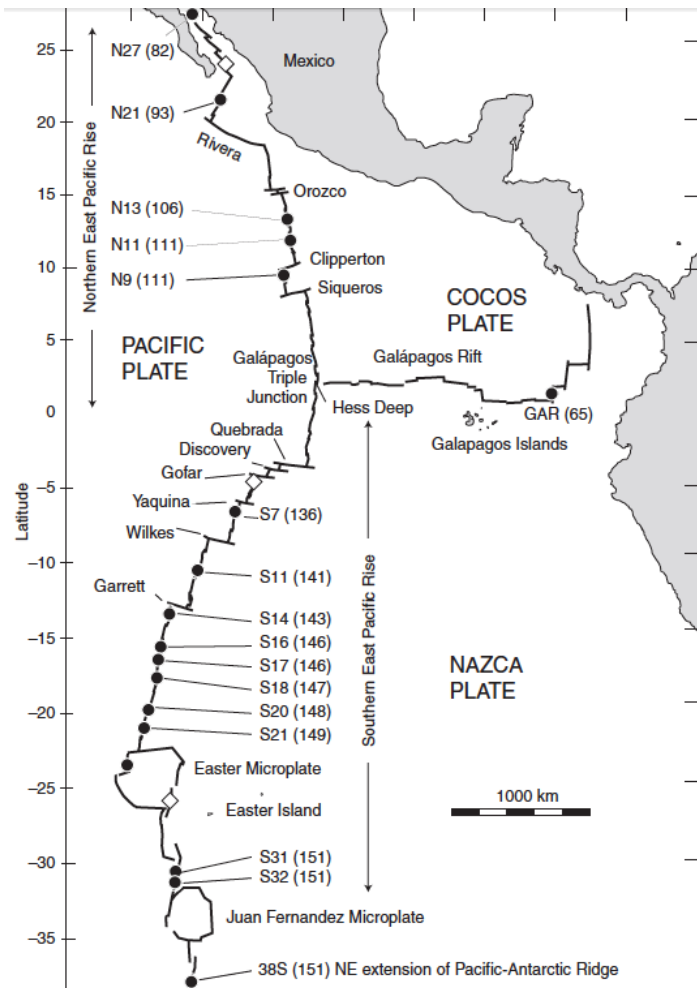
Teixeira, Serrao, Arnaud-Haond 2012. PloS ONE

Teixeira, Cambon-Bonavita, Serrao, Desbruyères, Arnaud-Haond 2010. J. Biogeography



Genetic differences between populations

Audzijonyte & Vrijenhoek 2010. Evolution
 Vrijenhoek 2010 Molecular Ecology



Patterns of differentiation in vent species from the northern (NEPR) and southern (SEPR) East Pacific Rise and Galapagos Rift (grey box). Figure is modified and updated from Audzijonyte & Vrijenhoek (2010), who estimated the correlations (IBD r)

Disturbances to MGR of the High Seas

Direct destruction of habitats and diversity by extractive activities (major impacts of local activities):

fisheries - target species, by-catch, habitat and ecosystem effects
mining – major impacts as technologies progress

Indirect destruction by global human pollution:

acidification

climate change and its effects (water stratification, land inputs)

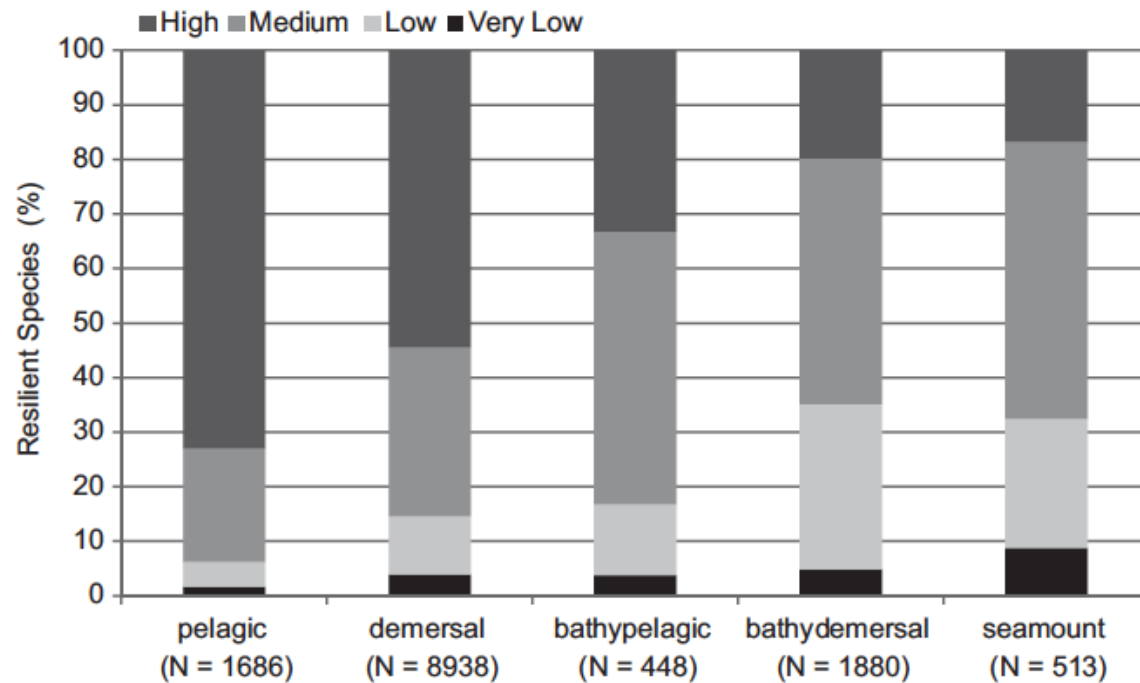
pollution convergence zones

Minimal or no impact:

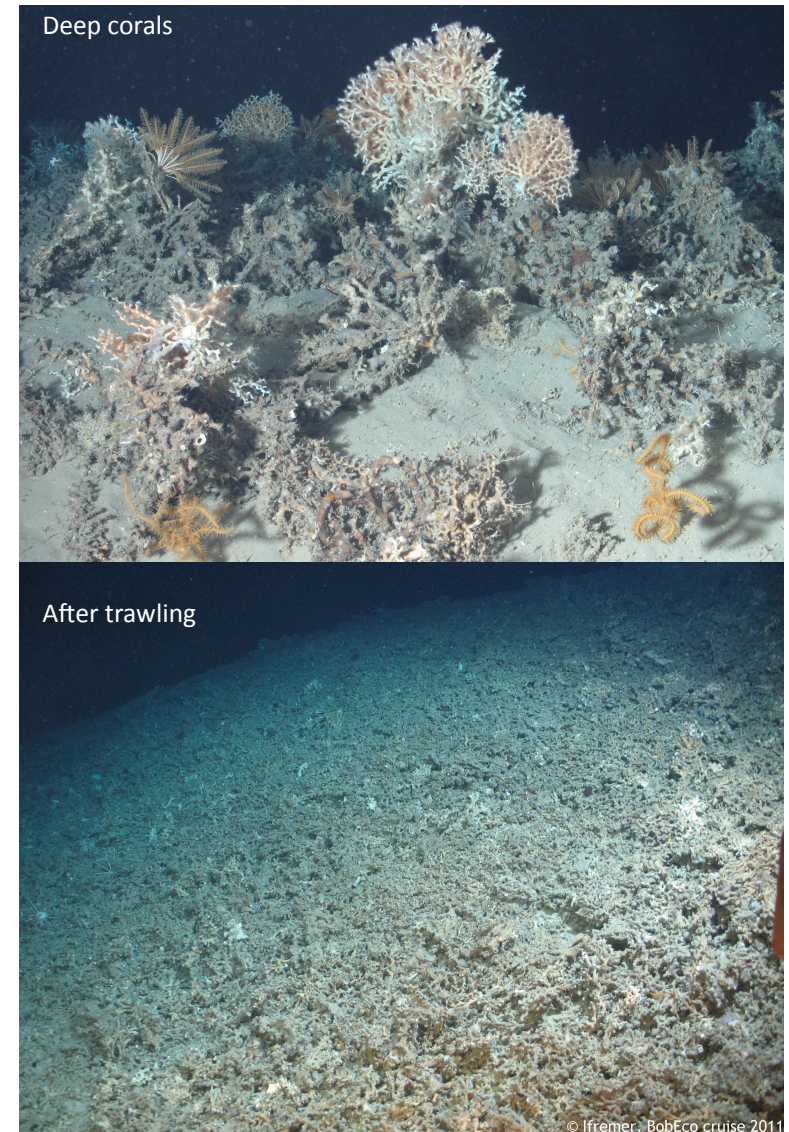
Genetic studies, bioprospecting - much information from minimal amount of sampling, below natural mortality rates.

Impacts on MGR

Seamount genetic resources especially vulnerable due to long, slow demographic processes



Norse, Brooke, Cheung, Clark, Ekeland, Froese, Gjerde, Haedrich, Heppell, Morato, Morgan, Pauly, Sumaila, Watson. Sustainability of deep-sea fisheries. Marine Policy. 2012.



necessary to discover and describe the
MGR of today and to share this knowledge
genetic loss happens unnoticed

Canyon du Croizic

the present/past natural state must be
known - shifting genetic baselines

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